

Figure 1. Phage-display selects out the OSCP. Semiquantitative PCR using primers specific for the OSCP sequence reveals enrichment of clones encoding the OSCP with each round of selection using immobilized Bz-423.

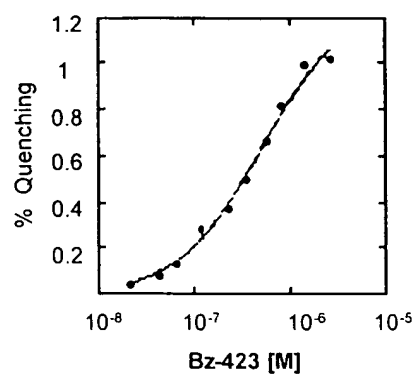


Figure 2. Binding isotherm of Bz-423 and purified human OSCP .

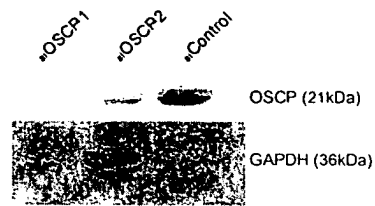


Figure 3. siRNA reduces the OSCP. 10^6 HEK 293 cells transfected with 0.84 μ g of siRNA duplexes specific for OSCP or scrambled control. After 72 h, protein expression determined by immunoblotting whole cell extracts. OSCP is reduced and GAPDH increased as glycolytic response.

5

10

15

FIGURE 4A.

	NCBI Accession Number	FOLD INCREASE
gene		
glyceraldehyde-3-phosphate dehydrogenase	M33197	23.02
ribosomal protein L28	NM_000892	21.62
heat shock 90kD protein 1, beta	A1216218	21.27
lysosomal-associated multispanning membrane protein-5	NM_008782	20.19
ferritin, light polypeptide	BG538564	19.50
Consensus includes gb:AJ249377.1 /DEF=Homo sapiens partial mRNA for human Ig lambda light chain variable region, clone MB91 (331 bp). /FEA=mRNA /GEN=IGLV /PROI AJ249377		18.95
Consensus includes gb:D84143.1 /DEF=Human Immunoglobulin (mAb59) light chain V region mRNA, partial sequence. /FEA=mRNA /PROD=immunoglobulin light chain V-J n D84143		18.76
gb:AJ25092.1 /DEF=Homo sapiens mRNA for single-chain antibody, complete cds. /FEA=CDS /PROD=immunoglobulin /DB_XREF=gi:3090425 /UG=Hs.249245 Homo sapiens	AJ25092	18.62
ribosomal protein L28	BF683426	18.53
Consensus includes gb:L48784 /DEF=Homo sapiens cDNA /FEA=mRNA /DB_XREF=gi:1068715 /UG=Hs.182426 ribosomal protein S2		17.74
profilin 1	L48784	17.74
ornithine decarboxylase antizyme 1	NM_005022	17.74
serine hydroxymethyltransferase 2 (mitochondrial)	AF090094	16.18
chaperonin containing TCP1, subunit 7 (eta)	NM_005412	15.75
ribosomal protein L8	NM_006429	15.12
macrophage migration inhibitory factor (glycosylation-inhibiting factor)	NM_000973	14.94
Consensus includes gb:D84140.1 /DEF=Human Immunoglobulin (mAb56) light chain V region mRNA, partial sequence. /FEA=mRNA /PROD=immunoglobulin light chain V-J n D84140		14.75
transketolase (Wernicke-Korsakoff syndrome)	NM_002415	14.48
ribosomal protein L18	BF696840	14.41
transketolase (Wernicke-Korsakoff syndrome)	NM_000979	14.38
ubiquitin-conjugating enzyme E21 (UBC9 homolog, yeast)	L12711	13.75
glyceraldehyde-3-phosphate dehydrogenase	AA910614	12.75
glyceraldehyde-3-phosphate dehydrogenase	M33197	12.69
nuclease sensitive element binding protein 1	BF689355	12.67
neutrophil cytosolic factor 4 (40kD)	BE966374	12.57
T-cell leukemia/lymphoma 1A	NM_013416	12.52
ribosomal protein L16a	X82240	12.46
glyceraldehyde-3-phosphate dehydrogenase	NM_000980	12.33
gb:L07950.1 /DEF=Homo sapiens MHC class I HLA B71 mRNA, complete cds. /FEA=CDS /GEN=HLA-B /PROD=MHC HLA B71 /DB_XREF=gi:307236 /FL=gb.L07950.1		12.29
ubiquitin carrier protein	BE561479	12.26
actin, beta	L07950	12.12
enolase 1, (alpha)	NM_014501	11.85
replication factor C (activator 1) 2 (40kD)	X00351	11.53
phosphoserine aminotransferase	U88968	11.48
MCM7 minichromosome maintenance deficient 7 (S. cerevisiae)	M87338	10.95
ribosomal protein S5	NM_021154	10.74
RNA polymerase II transcriptional regulation mediator (Med6, S. cerevisiae, homolog of)	AF279900	10.69
eukaryotic translation initiation factor 3, subunit 8 (110kD)	NM_001009	10.68
chaperonin containing TCP1, subunit 2 (beta)	NM_005466	10.65
CGH135 protein	BC000533	10.52
polymerase (DNA directed), delta 2, regulatory subunit (50kD)	AL545982	10.35
ribosomal protein L13	NM_016068	10.25
procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), beta polypeptide (protein disulfide isomerase; thyroid hormone binding protein p55)	NM_006230	9.93
HMT1 hnRNP methyltransferase-like 2 (S. cerevisiae)	AW574664	9.85
cold inducible RNA binding protein	J02763	9.83
ATP synthase, H+ transporting, mitochondrial F1 complex, delta subunit	NM_001536	9.86
pinin, desmosome associated protein	NM_001280	9.57
pyruvate kinase, muscle	BE798517	9.50
fusion, derived from t(12;16) malignant liposarcoma	AF112222	9.43
immunoglobulin heavy constant mu	NM_002654	9.40
maternal G10 transcript	L23516	9.42
major histocompatibility complex, class I, B	NM_004960	9.37
small EDRK-rich factor 2	U80139	9.32
heat shock 90kD protein 1, beta	NM_003910	9.30
Consensus includes gb:AF254822 /DEF=Homo sapiens SMARCA4 isoform (SMARCA4) gene, complete cds, alternatively spliced /FEA=CDS_2 /DB_XREF=gi:10946127 /UG=		9.29
protein phosphatase 1G (formerly 2C), magnesium-dependent, gamma isoform	D83043	9.29
glucose phosphate isomerase	NM_005770	9.05
proteasome (prosome, macropain) 26S subunit, non-ATPase, 8	AF275719	8.88
	AF254822	8.77
	NM_002707	8.76
	NM_000175	8.74
	NM_002812	8.74

ribosomal protein S19	BE259729	8.68
eukaryotic translation elongation factor 1 alpha 1	AL035687	8.48
gb.L23518.1 /DEF=Human Ig rearranged gamma-chain, V-DXP1-JH4b, complete cds. /FEA=mRNA /DB_XREF=gi:385220 /FL=gb:L23518.1	L23518	8.41
ATP synthase, H+ transporting, mitochondrial F1 complex, delta subunit	NM_001687	8.41
procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), beta polypeptide (protein disulfide isomerase; thyroid hormone binding protein p55)	NM_000918	8.38
endothelial differentiation-related factor 1	AB002282	8.37
acid phosphatase 1, soluble	BG035989	8.33
chaperonin containing TCP1, subunit 3 (alpha)	NM_005988	8.32
proteasome (prosome, macropain) subunit, alpha type, 7	AL078633	8.23
Consensus includes gb:AF043584.1 /DEF=Homo sapiens clone ASMmeg1-b1 Immunoglobulin lambda chain VJ region, (IGL) mRNA, partial cds. /FEA=mRNA /GEN=IGL /PRO AF043584	AF043584	8.21
non-POU-domain-containing, octamer-binding	L14599	8.09
CGI-51 protein	NM_015380	8.07
adaptor-related protein complex 2, mu 1 subunit	NM_004068	7.95
interferon regulatory factor 4	NM_002460	7.90
ribosomal protein S19	NM_001022	7.90
interferon stimulated gene (20kD)	NM_002201	7.88
dolichyl-diphosphooligosaccharide-protein glycosyltransferase	D28643	7.80
actin, beta	X00351	7.72
major histocompatibility complex, class II, DR alpha	M60333	7.69
major histocompatibility complex, class I, B	L42024	7.62
major histocompatibility complex, class II, DR beta 5	AJ287586	7.57
actin, gamma 1	AL567820	7.49
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (9kD, B9)	NM_004542	7.48
HLA-B associated transcript 1	NM_004640	7.47
ribosomal protein L18a	AC004692	7.36
H2A histone family, member O	A0313324	7.31
gb:U62824.1 /DEF=Homo sapiens HLA class I heavy chain (HLA-Cw*1701) mRNA, complete cds. /FEA=CDS /GEN=HLA-Cw*1701 /PROD=HLA class I heavy chain /DB_XREF U62824	U62824	7.31
major histocompatibility complex, class II, DR alpha	M60334	7.27
ribosomal protein S15	NM_001018	7.26
eukaryotic translation initiation factor 3, subunit 8 (110kD)	NM_003752	7.25
U8 snRNA-associated Sm-like protein LSM7	NM_016199	7.25
eukaryotic translation elongation factor 2	NM_001961	7.22
membrane-spanning 4-domains, subfamily A, member 1	X12530	7.13
eukaryotic translation initiation factor 3, subunit 8 (110kD)	AA678705	7.11
spermidine synthase	NM_003132	7.09
polymerase (RNA) II (DNA directed) polypeptide J (13.3kD)	BG335629	7.09
ribosomal protein S2	NM_002952	7.03
chemokine (C-X-C motif), receptor 4 (fusin)	AF348491	7.03
similar to HYPOTHETICAL 34.0 KDA PROTEIN ZK795.3 IN CHROMOSOME IV	BC003629	7.00
DEAD-box protein abstrakt	BE747342	6.98
immunoglobulin heavy constant mu	AF081459	6.94
DKFZP564M182 protein	NM_016222	6.93
nuclear RNA helicase, DECD variant of DEAD box family	BC001872	6.90
stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing protein)	AK000822	6.88
immunoglobulin heavy constant mu	NM_005804	6.88
muS homolog 6 (E. coli)	BE866560	6.80
tubulin, beta 4	BG340548	6.76
major histocompatibility complex, class II, DR beta 1	D89846	6.75
T cell receptor beta locus	U65565	6.73
major histocompatibility complex, class II, DP alpha 1	M15564	6.72
ribonucleotide reductase M1 polypeptide	M27487	6.62
signal sequence receptor, delta (translocon-associated protein delta)	A1692874	6.62
seryl-RNA synthetase	NM_006280	6.59
eukaryotic translation initiation factor 2, subunit 3 (gamma, 52kD)	NM_006513	6.55
protein phosphatase 1, regulatory subunit 7	NM_001415	6.54
ancient ubiquitous protein 1	BF718769	6.52
glutathione S-transferase pi	NM_012103	6.52
polymerase (RNA) III (DNA directed) polypeptide K (12.3 kDa)	NM_000852	6.49
ubiquinol-cytochrome c reductase core protein 1	NM_016310	6.42
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	NM_003365	6.42
splicing factor, arginine/serine-rich 5	AB033605	6.41
	AW084582	6.40

SKB1 homolog (S. pombe)
 cell death-regulatory protein GRIM19
 U6 snRNA-associated Sm-like protein
 eukaryotic translation initiation factor 3, subunit 2 (beta, 36kD)
 mitochondrial ribosomal protein S2
 D123 gene product
 replication factor C (activator 1) 2 (40kD)
 membrane-spanning 4-domains, subfamily A, member 1
 Rab acceptor 1 (prenylated)
 eukaryotic translation initiation factor 3, subunit 9 (eta, 116kD)
 ubiquitin C
 valosin-containing protein
 ATP synthase, H⁺ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 1
 CD22 antigen
 B lymphoid tyrosine kinase
 mitochondrial ribosomal protein S34
 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASH1)
 mitochondrial ribosomal protein L9
 ribosomal protein S2
 actin binding LIM protein 1
 syntrophin 16
 major histocompatibility complex, class II, DR beta 4
 protein kinase, DNA-activated, catalytic polypeptide
 ribosomal protein S28
 eukaryotic translation initiation factor 4A, isoform 1
 ribosomal protein, large, PO
 NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae)
 ATP citrate lyase
 Consensus includes gb:BF9794.19 /FEA=EST /DB_XREF=gi:12346634 /DB_XREF=est:602286246F1 /CLONE=IMAGE-4373914 /UG=Hs.119122 ribosomal protein L13a
 glutamyl-tRNA synthetase
 Consensus includes gb:AF005487.1 /DEF=Homo sapiens MHC class II antigen (DRB6) mRNA, HLA-DRB6*0201 allele, sequence. /FEA=mrna /DB_XREF=gi:5915883 /UG=H
 eukaryotic translation initiation factor 3, subunit 9 (eta, 116kD)
 polymerase (RNA) II (DNA directed) polypeptide L (7.6kD)
 mitochondrial ribosomal protein S12
 ATP synthase, H⁺ transporting, mitochondrial F0 complex, subunit 2
 putative human HLA class II associated protein I
 hypothetical protein
 membrane component, chromosome 11, surface marker 1
 hypothetical protein PRO1847
 proteasome (prosome, macropain) subunit, beta type, 7
 cell division cycle 2-like 2
 similar to RIKEN cDNA 2310040G17 gene
 putative transmembrane protein; homolog of yeast Golgi membrane protein Yif1p (Yip1p-interacting factor)
 tylophanyl-RNA synthetase
 ribosomal protein, large, PO
 transcription factor Dp-1
 guanine nucleotide binding protein (G protein), beta polypeptide 1
 actin, gamma 1
 hypothetical protein R33729_1
 CDW52 antigen (CAMPATH-1 antigen)
 ribosomal protein, large, PO
 actin related protein 23 complex, subunit 2 (34 kD)
 guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1
 eukaryotic translation initiation factor 4A, isoform 1
 ubiquitin C
 polymerase (RNA) II (DNA directed) polypeptide J (13.3kD)
 hypothetical protein MGC4875
 ribosomal protein L10
 v-yes-1 Yarnaguchi sarcoma viral related oncogene homolog
 calpain, small subunit 1
 lymphocyte-specific protein tyrosine kinase
 anaphase promoting complex subunit 5
 ubiquitin-conjugating enzyme E2G 2 (UBC7 homolog, yeast)

NM_006109 6.32
 NM_015965 6.20
 NM_012321 6.18
 U38764 6.17
 NM_016034 6.16
 NM_006023 6.15
 NM_002914 6.09
 BC002807 6.02
 NM_006423 5.97
 BC001173 5.97
 M28880 5.94
 AF100752 5.93
 AL080089 5.93
 X52785 5.89
 NM_001715 5.88
 NM_023936 5.88
 NM_005004 5.86
 AB049636 5.85
 A183766 5.84
 NM_006720 5.84
 AK026970 5.83
 NM_021863 5.82
 U34994 5.81
 AC005011 5.80
 NM_001416 5.76
 BC003655 5.75
 AF155235 5.75
 U18197 5.74
 BF979419 5.74
 NM_005051 5.73
 NM_003751 5.70
 BC005803 5.66
 NM_021107 5.66
 NM_004889 5.65
 BE560202 5.64
 NM_016459 5.58
 BG258784 5.57
 AF119855 5.57
 NM_002799 5.57
 AF067524 5.57
 BF972185 5.57
 NM_020470 5.56
 M61715 5.56
 NM_001002 5.56
 NM_007111 5.55
 AU141124 5.54
 AU145192 5.53
 AC005339 5.53
 NM_001803 5.52
 A953822 5.51
 AF279893 5.50
 NM_006098 5.49
 BC006210 5.47
 AB008010 5.47
 AW402635 5.40
 AL118502 5.39
 NM_006013 5.35
 M79321 5.33
 AD001527 5.31
 NM_005356 5.31
 BC001081 5.29
 BG395660 5.29

kinesin 2 (60-70kD)	AA284075	5.29
hematological and neurological expressed 1	NM_018185	5.27
ribosomal protein, large, P0	BC005863	5.26
immunoglobulin heavy constant mu	S74839	5.25
proteasome (prosome, macropain) subunit, beta type, 8 (large multifunctional protease 7)	U17496	5.24
ribosomal protein S3	U14990	5.22
proteasome (prosome, macropain) 26S subunit, ATPase, 3	AL545523	5.21
coronin, actin binding protein, 1A	U34690	5.19
nuclear distribution gene C (A. nidulans) homolog	AF241788	5.17
neutrophil cytosolic factor 4 (40kD)	NM_000631	5.17
gb:M24688.1 /DEF=Human Ig rearranged H-chain V-region mRNA (C-D-JH4), complete cds. /FEA=mRNA /GEN=IGH@_DB_XREF=gi:185198 /FL=gb:M24688.1	M24688	5.16
splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate splicing factor)	NM_006924	5.16
Ov1 antigen	M12879	5.11
aristae homolog 2 (Drosophila)	BC004422	5.09
bromodomain-containing 2	D42040	5.09
major histocompatibility complex, class I, F	AW514210	5.08
succinate dehydrogenase complex, subunit B, iron sulfur (lp)	NM_003000	5.08
NADH dehydrogenase (ubiquinone) Fe-S protein 3 (30kD) (NADH-coenzyme Q reductase)	NM_004551	5.07
farnesyl-diphosphate farnesyltransferase 1	BC003573	5.01
transgelin 2	NM_003584	4.98
cytochrome c oxidase subunit IV isoform 1	NM_001861	4.98
integrin beta 4 binding protein	AF022229	4.95
HIF-1 responsive RTP801	NM_019058	4.93
hypothetical protein PRO1068	NM_018573	4.90
putative breast adenocarcinoma marker (32kD)	NM_014453	4.88
protein tyrosine phosphatase, receptor type, O	U20489	4.88
hypothetical protein DKFZp434N185	NM_025205	4.86
tubulin alpha 6	BC005946	4.85
ribosomal protein L13	AA789278	4.85
Lysosomal-associated multispanning membrane protein-5	AI589086	4.83
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1)	M18468	4.83
ribosomal protein, large P2	NM_001004	4.82
eukaryotic translation initiation factor 4 gamma, 1	AF104913	4.79
transcription elongation factor 8 (SIII), polypeptide 2 (18kD, elongin B)	NM_007108	4.78
karyopherin (importin) beta 3	NM_002271	4.76
lymphocyte cytosolic protein 1 (L-plastin)	J02923	4.74
peroxiredoxin 1	L19184	4.70
heat shock protein 75	NM_016282	4.70
HLA-G histocompatibility antigen, class I, G	AF228990	4.70
ribosomal protein S14	AF118710	4.69
cullin 1	NM_003592	4.69
heat shock 70kD protein 4	BC002526	4.68
NADH dehydrogenase (ubiquinone) flavoprotein 1 (51kD)	AF082131	4.65
HSPC274 protein	NM_014145	4.63
Consensus includes gb:BE305165 /FEA=EST /DB_XREF=gi:9177184 /DB_XREF=est601186685T1 /CLONE=IMAGE:2959580 /UG=Hs.100623 phospholipase C, beta 3, neigl	BE305165	4.60
Consensus includes gb:BE305165 /FEA=EST /DB_XREF=gi:9177184 /DB_XREF=est601186685T1 /CLONE=IMAGE:2959580 /UG=Hs.100623 phospholipase C, beta 3, neigl	D55716	4.60
MCM7 minichromosome maintenance deficient 7 (S. cerevisiae)	NM_003406	4.56
lysine 3-monooxygenase/hypophan 5-monooxygenase activation protein, zeta polypeptide	U10485	4.55
lymphoid-restricted membrane protein	BG389015	4.53
tumor protein D52	J04564	4.53
small nuclear ribonucleoprotein polypeptides B and B1	U38979	4.52
postmeiotic segregation increased 2-like 9	NM_002466	4.51
v-myb myeloblastosis viral oncogene homolog (avian)-like 2	AF043179	4.46
T cell receptor beta locus	AW150953	4.45
7-dehydrocholesterol reductase	AL523776	4.45
hypothetical protein FLJ20113	BC000733	4.44
eukaryotic translation initiation factor 3, subunit 4 (delta, 44kD)	AB028869	4.41
baculoviral IAP repeat-containing 3 (survivin)	NM_006152	4.39
lymphoid-restricted membrane protein	M80281	4.39
APEX nuclease (multifunctional DNA repair enzyme)	NM_028202	4.39
Consensus includes gb:Z82202 /DEF=Human DNA sequence from clone RP1-34P24 on chromosome 22 Contains a pseudogene similar to ribosomal protein L35, ESTs, STs	NM_018694	4.38
HSV1 binding protein	AA112507	4.38
U6 snRNA-associated Sm-like protein	AL559122	4.38
T cell receptor beta locus		

casein kinase 2, beta polypeptide	NM_001320	4.38
putative methyltransferase	NM_017528	4.37
VPS28 protein	NM_016208	4.35
lysosomal-associated membrane protein 1	NM_005561	4.33
bone marrow stromal cell antigen 2	NM_004335	4.32
ribosomal protein L13a	BF942308	4.32
transmembrane protein 4	BC001027	4.32
calreticulin	AD000092	4.30
nuclear RNA export factor 1	BC004904	4.30
polymerase (RNA) II (DNA directed) polypeptide E (25kD)	AI554759	4.28
nucleosome assembly protein 1-like 4	NM_005969	4.25
gb:AY014272.1 /DEF=Homo sapiens FKSG30 (FKSG30) mRNA, complete cds. /FEA=mRNA (GEN=FKSG30 /PROD=FKSG30 /DB_XREF=gi:12408251 /UG=Hs.315492 Homc	NM_007773	4.25
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)	AI768378	4.24
KIA00818 gene product	NM_005574	4.23
LIM domain only 2 (homobin-like 1)	BC003129	4.22
non-POU-domain-containing, octamer-binding	NM_005729	4.19
peptidylprolyl isomerase F (cyclophilin F)	BE671379	4.18
zinc finger protein 207	AV702405	4.18
enopamil binding protein (sterol isomerase)	NM_016209	4.17
unknown	NM_006118	4.17
H31 binding protein	NM_001098	4.16
aconitase 2, mitochondrial	NM_005324	4.15
H3 histone, family 3B (H3.3B)	U72514	4.14
C2f protein	BC001675	4.14
ribosomal protein L13a	NM_002945	4.12
replication protein A1 (70kD)	NM_002808	4.12
prolaseome (prosome, macropain) 26S subunit, non-ATPase, 2	NM_001550	4.11
gp:AY014272.1 /DEF=Homo sapiens FKSG30 (FKSG30) mRNA, complete cds. /FEA=mRNA (GEN=FKSG30 /PROD=FKSG30 /DB_XREF=gi:12408251 /UG=Hs.315492 Homc	M90684	4.11
interferon-related developmental regulator 1	NM_016337	4.11
HLA-G histocompatibility antigen, class I, G	U23946	4.10
RNB6	NM_003145	4.09
RNA binding motif protein 5	NM_005273	4.09
signal sequence receptor, beta (translocon-associated protein beta)	NM_001029	4.08
guanine nucleotide binding protein (G protein), beta polypeptide 2	M12423	4.07
ribosomal protein S26	NM_012458	4.07
T cell receptor alpha locus	NM_004111	4.06
translocase of inner mitochondrial membrane 13 homolog B (yeast)	NM_014294	4.06
flap structure-specific endonuclease 1	BC003381	4.06
translocating chain-associating membrane protein	NM_012456	4.05
KIAA0217 protein	NM_005919	4.04
translocase of inner mitochondrial membrane 10 homolog (yeast)	BC004949	4.03
MADS box transcription enhancer factor 2, polypeptide B (myocyte enhancer factor 2B)	M31183	4.03
tubulin alpha 6	AF217197	4.02
major histocompatibility complex, class I, E	BC005147	4.02
fuse-binding protein-interacting repressor	AI987323	4.01
ATP synthase, H+ transporting, mitochondrial F1 complex, alpha subunit, isoform 1, cardiac muscle	AF321125	4.00
DNA replication factor	BG028844	4.00
HLA-B associated transcript 3	NM_005001	4.00
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5kD, B14.5a)	U74223	4.00
poly(rC) binding protein 1	NM_005724	3.99
tetraspan 3	AI972144	3.99
hydroxycy-Coenzyme A dehydrogenase/3-ketocyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit	NM_000075	3.98
cyclin-dependent kinase 4	BE560461	3.98
solute carrier family 2 (facilitated glucose/fructose transporter), member 5	BG257762	3.98
hypothetical protein	M15565	3.96
T cell receptor alpha locus	NM_006409	3.95
actin related protein 2/3 complex, subunit 1A (41 kD)	L34703	3.94
T cell receptor alpha locus	AW001847	3.94
amyloid beta (A4) precursor-like protein 2	NM_006327	3.93
translocase of inner mitochondrial membrane 23 homolog (yeast)	AL535380	3.93
B-cell translocation gene 1, anti-proliferative	BC003049	3.92
PAI-1 mRNA-binding protein	U39945	3.92
adenyate kinase 2		

cytochrome c oxidase subunit VIc	A3382702	3.9
TAP binding protein (tapasin)	AF029750	3.91
ribosomal protein L27a	NM_000990	3.91
ribosomal protein L27a	NM_002708	3.91
protein phosphatase 1, catalytic subunit, alpha isoform	BG531983	3.90
Consensus includes gb:RG531983 /FEA=EST /DB_XREF=g :3523521 /DB_XREF=est:602561007F1 /CLONE=IMAGE:4699176 /UG=Hs.108104 ubiquitin-conjugating enzyme	BG531983	3.90
Consensus includes gb:RG334495 /FEA=EST /DB_XREF=g :31140933 /DB_XREF=est:602461128F1 /CLONE=IMAGE:4577718 /UG=Hs.228201 ESTs. Moderately similar to / BG334495	BG334495	3.90
leucine rich repeat (in FLII) interacting protein 1	NM_004735	3.89
ribosomal protein, large, P1	NM_001003	3.89
cold shock domain protein A	NM_003651	3.89
Tu translation elongation factor, mitochondrial	NM_003321	3.89
ribosomal protein S18	NM_022551	3.86
KIAA0748 protein	A4522514	3.86
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform	NM_014225	3.85
ubiquitin-activating enzyme E1 (A1SST and BN75 temperature sensitivity complementing)	NM_003334	3.85
mitochondrial ribosomal protein S16	NM_016065	3.84
8-phosphogluconolactonase	NM_012088	3.84
CCAAT/enhancer binding protein (C/EBP), gamma	NM_001806	3.83
major histocompatibility complex, class I, E	NM_005516	3.82
coatomer protein complex, subunit epsilon	NM_007263	3.81
chloride intracellular channel 1	AF034607	3.80
hypothetical protein FLJ20340	NM_011773	3.80
CD27-binding (Siva) protein	AF033111	3.80
postmitotic segregation increased 2-like 11	U38980	3.79
Consensus includes gb:X02189 /DEF=F.H.sapiens adenosine deaminase (ADA) gene 5 flanking region and exon 1 (and joined CDS) /FEA=CDS /DB_XREF=g :26358 /UG=Hs.1	X02189	3.79
hypothetical protein	A1670847	3.78
Consensus includes gb:BF530535 /FEA=EST /DB_XREF=g :11617898 /DB_XREF=est:602071788F1 /CLONE=IMAGE:4214660 /UG=Hs.77865 KIAA0102 gene product /FL=g	BF530535	3.78
FK506 binding protein precursor	NM_006831	3.77
CDW52 antigen (CAMPATH-1 antigen)	N90868	3.76
tyrosine 3-monooxygenase/tyrosophan 5-monooxygenase activation protein, epsilon polypeptide	U28836	3.75
translocase of inner mitochondrial membrane 44 homolog (yeast)	NM_006156	3.75
neural precursor cell expressed, developmentally down-regulated 8	NM_006026	3.75
H1 histone family, member X	N28665	3.74
KIAA0618 gene product	NM_005826	3.74
heterogeneous nuclear ribonucleoprotein R	U51007	3.74
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	NM_002795	3.73
proteasome (prosome, macropain) subunit, beta type, 3	A142677	3.72
glutaryl-prolyl-PRNA synthetase	A1697055	3.71
GCH1 general control of amino-acid synthesis 1-like 1 (yeast)	NM_001404	3.71
eukaryotic translation elongation factor 1 gamma	AF113682	3.71
thymopoesin	AF113682	3.71
Consensus includes gb:BE731738 /FEA=EST /DB_XREF=g :10145730 /DB_XREF=est:601568154F1 /CLONE=IMAGE:3842844 /UG=Hs.192337 peptidylprolyl isomerase A (c	BE731738	3.71
cytochrome c oxidase subunit Vb	NM_001862	3.70
mitochondrial carrier homolog 2	NM_014342	3.68
tyrosine 3-monooxygenase/tyrosophan 5-monooxygenase activation protein, zeta polypeptide	U28964	3.68
interleukin enhancer binding factor 3, 90kD	AF147209	3.68
enhancer of invasion 10	NM_021178	3.67
RuvB-like 2 (E. coli)	NM_006866	3.65
small nuclear ribonucleoprotein polypeptide A	NM_004596	3.65
hypothetical protein FLJ22837	NM_025165	3.65
myosin 1D	AA621962	3.64
cytochrome P450, subfamily 51 pseudogene 2	U40053	3.64
SNRPN upstream reading frame	NM_022804	3.63
ribosomal protein L13a	NM_012423	3.63
H2B histone family, member G	NM_003522	3.63
stress-induced-phosphoprotein 1 (Hsp70Hsp90-organizing protein)	AL553320	3.63
leucine-rich repeat protein, neuronal 1	A1654857	3.63
nuclear distribution gene C (A.nidulans) homolog	NM_006600	3.63
hydroxyacyl-Coenzyme A dehydrogenase, type II	NM_004493	3.62
MCN5 minichromosome maintenance deficient 5, cell division cycle 46 (S. cerevisiae)	NM_006739	3.62
PTD008 protein	NM_016145	3.61
eukaryotic translation elongation factor 1 alpha 1	NM_001402	3.60
adenine phosphoribosyltransferase	AA927724	3.60
GDP-mannose 4,6-dehydratase	NM_001500	3.59

PRP9 pre-mRNA processing factor 8 homolog (yeast)
 hepatitis delta antigen-interacting protein A
 Consensus includes gb:Z95126 /DEF=Human DNA sequence from PAC 30P20 on chromosome Xq21.1-Xq21.3. Contains set pseudogene, ESTs and STS /FEA=COS /DB_XR_Z95126
 Williams-Beuren syndrome chromosome region 5
 guanylate kinase 1
 intercellular adhesion molecule 2
 nucleolar protein family A, member 3 (HACA small nucleolar RNPs)
 G protein-coupled receptor 18
 major histocompatibility complex, class II, DQ alpha 1
 ribosomal protein L15
 ATP-binding cassette, sub-family F (GCM20), member 2
 KIAA0618 gene product
 BCL2-interacting killer (apoptosis-inducing)
 adaptor-related protein complex 2, sigma 1 subunit
 Consensus includes gb:BF695847 /FEA=EST /DB_XREF=gi:11981255 /DB_XREF=est:601852205F1 /CLONE=IMAGE:4076232 /UG=Hs.301917 YDD19 protein
 ARP2 actin-related protein 2 homolog (yeast)
 structure specific recognition protein 1
 myosin, light polypeptide 6, alkali, smooth muscle and non-muscle
 delta sleep inducing peptide, immunoreactor
 Consensus includes gb:A001784 /FEA=EST /DB_XREF=gi:3202255 /DB_XREF=est:rt41g06.s1 /CLONE=IMAGE:1619386 /UG=Hs.308332 ESTs, Highly similar to A42735 rb A101784
 mercaptopyruvate sulfurtransferase
 GNAS complex locus
 hypothetical protein FLJ10359
 DKFZp434J1813 protein
 lymphotoxin beta (TNF superfamily, member 3)
 Consensus includes gb:AJ011414.1 /DEF=Homo sapiens mRNA for plexin-B1 plasma membrane receptor, truncated splice variant (plexin-B1SEP gene). /FEA=mRNA /GEN=AJ011414
 DKFZP586M1523 protein
 cytochrome c oxidase subunit Vb
 ribonuclease H1
 methionine-tRNA synthetase
 immunoglobulin lambda locus
 proteasome (prosome, macropain) 26S subunit, non-ATPase, 3
 myosin, light polypeptide 6, alkali, smooth muscle and non-muscle
 insulin induced gene 1
 ATP synthase, H+ transporting, mitochondrial F1 complex, beta polypeptide
 thyroid autoantigen 70kD (Ku antigen)
 Consensus includes gb:AK027225.1 /DEF=Homo sapiens cDNA: FLJ23572 fls, clone LNG12403. /FEA=mRNA /DB_XREF=gi:10440306 /UG=Hs.201593 Homo sapiens cDNA: AK027225
 branched chain aminotransferase 1, cytosolic
 FK506 binding protein 1A (12kD)
 Huntingtin interacting protein B
 chromatin-specific transcription elongation factor, 140 kDa subunit
 similar to mouse Gli3 or D. melanogaster transcription factor IIB
 zinc finger protein 313
 POP7 (processing of precursor, S. cerevisiae) homolog
 structure specific recognition protein 1
 mitochondrial ribosomal protein L23
 isopentenyl-diphosphate delta isomerase
 Xq28, 2000bp sequence contig, ORF
 glutathione peroxidase 1
 hypothetical protein MGC5585
 GNAS complex locus
 cytosolic acyl coenzyme A thioester hydrolase
 phosphoinositide-3-kinase, regulatory subunit, polypeptide 3 (p55, gamma)
 A kinase (PRKA) anchor protein 2
 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2 (8kD, AGGG)
 serine hydroxymethyltransferase 2 (mitochondrial)
 malic enzyme 2, NAD(+)-dependent, mitochondrial
 oxysterol binding protein-like 2
 HSPC142 protein
 RNA binding motif protein 10
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4
 DNA segment on chromosome 19 (unique) 1177 expressed sequence
 threonine-tRNA synthetase

NM_008445 3.59
 NM_008848 3.59
 AF257135 3.58
 BC008249 3.57
 NM_008873 3.57
 NM_018648 3.57
 AF261135 3.55
 BC397856 3.54
 AF279803 3.54
 NM_005692 3.54
 AA514822 3.53
 NM_001197 3.53
 NM_021575 3.53
 BF695847 3.52
 NM_005722 3.51
 NM_003146 3.51
 AA419227 3.51
 AL110191 3.50
 NM_001784 3.49
 NM_021126 3.45
 AF064082 3.45
 NM_018072 3.45
 BG168666 3.45
 NM_002341 3.44
 NM_001414 3.44
 BF063898 3.43
 BC008229 3.41
 NM_002836 3.41
 NM_004990 3.40
 XG3006 3.40
 NM_002809 3.40
 BE734356 3.39
 BE300521 3.37
 NM_001686 3.37
 NM_001469 3.36
 NM_007225 3.35
 NM_005604 3.35
 BC001002 3.34
 AF049103 3.33
 NM_007192 3.33
 NM_013242 3.33
 AL031685 3.32
 BC001430 3.32
 BE795648 3.31
 AL832239 3.31
 NM_004508 3.30
 BE076218 3.29
 NM_000581 3.29
 NM_024057 3.28
 AF088184 3.27
 NM_007274 3.27
 BE622627 3.27
 BE679367 3.27
 NM_004546 3.27
 AW180316 3.26
 MS5905 3.26
 BC000296 3.25
 BC006244 3.25
 AL137421 3.25
 A1744900 3.25
 NM_006114 3.25
 NM_003191 3.25

Consensus includes gb:X04802 /DEF=Homo sapiens UBBP2 pseudogene for ubiquitin UBB /FEA=CDS /DB_XREF=gi:37582 /UG=Hs.247890 Homo sapiens UBBP2 pseudoge	X04802	3.25
glyogenin	NM_004130	3.24
GNAS complex locus	NM_000516	3.24
ribosomal protein L10	AL031276	3.24
macrophage erythroblast attachr	NM_005882	3.24
Ewing sarcoma breakpoint region 1	BC006927	3.24
isodizate dehydrogenase 3 (NAD+) gamma	NM_004135	3.23
Consensus includes gb:AL118510 /DEF=Human DNA sequence from clone RPS-858M22 on chromosome 20 Contains a pseudogene similar to 40S ribosomal protein S10, ST	AL118510	3.23
paired immunoglobulin-like receptor beta	NM_013440	3.23
KIAA0729 protein	AW502434	3.22
serine palmitoyltransferase, long chain base subunit 2	U15555	3.22
endothelial differentiation-related factor 1	AB002282	3.22
kinesin-like 4	AC002301	3.21
ubiquitin B	X04801	3.21
siyltransferase 9 (CMP-NeuAc:lactosylceramide alpha-2,3-sialyltransferase; GM3 synthase)	NM_003896	3.21
eukaryotic translation elongation factor 1 alpha 1	NM_001403	3.20
chromosome 12 open reading frame 8	NM_006817	3.20
alanine-RNA synthetase	NM_001605	3.20
hypothetical protein FLJ11305	NM_018386	3.20
NADH dehydrogenase (ubiquinone) Fe-S protein 8 (23kD) (NADH-coenzyme Q reductase)	NM_002496	3.19
N-acylaminoacyl-peptide hydrolase	NM_001640	3.19
peptidylprolyl isomerase E (cyclophilin E)	AF042386	3.18
branched chain aminotransferase 1, cytosolic	AI652662	3.18
zinc finger protein 163 (RING finger, C3HC4 type)	BC000832	3.18
amino acid transporter 2	NM_018976	3.17
chromosome 14 open reading frame 3	NM_012111	3.17
SKI-interacting protein	AV689564	3.16
interferon regulatory factor 3	NM_001571	3.16
polyglutamine binding protein 1	AB041836	3.16
ubiquitin specific protease 7 (herpes virus-associated)	NM_003470	3.15
glutamic-pyruvate transaminase (alanine aminotransferase)	NM_005309	3.15
CD48 antigen (B-cell membrane protein)	NM_001778	3.15
golgi associated, gamma asadplin ear containing, ARF binding protein 2	BC000284	3.15
oxidase (Cytochrome c) assembly 1-like	BC001669	3.15
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	AI122754	3.14
RAM binding protein 1	NM_002882	3.14
mitochondrial solute carrier	BE677761	3.14
flightless I homolog (Drosophila)	AI830227	3.14
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 42kD	NM_001695	3.14
Consensus includes gb:BG537190 /FEA=EST /DB_XREF=gi:13528922 /DB_XREF=est:602565589F1 CLONE=IMAGE:4680079 /UG=Hs.111334 ferritin, light polypeptide	BG537190	3.13
CD27-binding (Sive) protein	NM_006427	3.13
FK506 binding protein precursor	NM_016594	3.13
succinate dehydrogenase complex, subunit A, flavoprotein (Fp)	NM_004168	3.12
chromosome condensation 1	NM_001269	3.11
PRKC, apoptosis, WT1, regulator	AI336206	3.11
villin 2 (ezrin)	J05021	3.11
glutamic-oxaloacetic transaminase 1, soluble (aspartate aminotransferase 1)	BC000498	3.11
splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila)	BC000451	3.11
hypothetical protein FLJ10869	NM_018292	3.10
interleukin 21 receptor	NM_021798	3.10
G8 protein	NM_016947	3.10
lysine 3-monoxygenase/hyptophan 5-monoxygenase activation protein, theta polypeptide	NM_008826	3.10
isopentenyl-diphosphate delta isomerase	BC005247	3.10
transforming growth factor, beta 1	BC000125	3.10
nicotinamide nucleotide transhydrogenase	NM_012343	3.10
homeo box A1	AC004079	3.10
CD37 antigen	NM_001774	3.10
G1 to S phase transition 1	NM_002094	3.09
ribosomal protein S8	NM_001012	3.09
glycoprotein, synaptic 2	NM_004668	3.09
suppression of tumorigenicity 14 (colon carcinoma, matriplase, epithin)	NM_021878	3.09
synaptophysin-like protein	AI768845	3.08
proteasome (prosome, macropain) 26S subunit, non-ATPase, 9	NM_002813	3.08

Fanconi anemia, complementation group A	AV083279	3.08
zinc finger protein 259	NM_003904	3.07
translin	AA435302	3.07
WD repeat domain 1	AB010427	3.07
ubiquitin-conjugating enzyme E2M (UBC12 homolog, yeast)	NM_003969	3.07
CCAAT/enhancer binding protein (C/EBP), beta	AL564683	3.07
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 1	NM_004939	3.06
NADH dehydrogenase (ubiquinone) Fe-S protein 8 (23kD) (NADH-coenzyme Q reductase)	NM_002496	3.06
immunoglobulin lambda joining 3	AV698647	3.06
H2A histone family, member E	NM_021066	3.05
gb:AF008937.1 DEF-Homo sapiens syntaxin-16C mRNA, complete cds. /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:2352817 /UG=Hs.102178 syntaxin 16 /FL=gb:AF008937	AF008937	3.05
block of proliferation 1	BC491842	3.05
interleukin enhancer binding factor 3, 90kD	AF141870	3.05
SMC4 structural maintenance of chromosomes 4-like 1 (yeast)	NM_009496	3.05
hypothetical protein FLJ12973	NM_024908	3.05
HLA-B associated transcript 3	BC003133	3.05
gb:Z25433.1 /DEF=H.sapiens protein-serine/threonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405744 /FL=gb:Z25433	Z25433	3.04
inhibitor of DNA binding 1, dominant negative helix-loop-helix protein	D13889	3.03
Consensus includes gb:AA680302 /FEA=EST /DB_XREF=gi:2656270 /DB_XREF=est:ae83405.s1 /CLONE=IMAGE:869193 /UG=Hs.181125 immunoglobulin lambda locus	AA680302	3.03
transcriptional regulator protein	NM_008710	3.03
COP9 homolog	AF081258	3.03
chromodomain protein, Y chromosome-like	J04984	3.03
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	BE262760	3.03
ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast)	NM_007273	3.03
B-cell associated protein	NM_002080	3.02
glutamic-oxaloacetic transaminase 2, mitochondrial (aspartate aminotransferase 2)	AA670344	3.02
villin 2 (ezrin)	NM_005258	3.02
GTP cyclohydrolase 1 feedback regulatory protein	AF119850	3.01
eukaryotic translation elongation factor 1 gamma	NM_016050	3.01
mitochondrial ribosomal protein L11	NM_001783	3.01
CD79A antigen (immunoglobulin-associated alpha)	BC001360	3.00
ras homolog gene family, member A	BC000779	3.00
KIAA1007 protein	NM_007318	2.99
presenilin 1 (Alzheimer disease 3)	BE551340	2.99
KIAA0974 protein	M87790	2.99
immunoglobulin lambda locus	NM_002716	2.98
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	BF340083	2.98
solute carrier family 1 (glutamate/neutral amino acid transporter), member 4	NM_002394	2.98
solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2	M65254	2.98
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	BC003005	2.97
inactive progesterone receptor, 23 kD	NM_008396	2.97
Spogen's syndrome/dermatoma autoantigen 1	NM_004639	2.97
HLA-B associated transcript 3	BF116254	2.97
triophosphate isomerase 1	S67779	2.96
B-cell CLL/lymphoma 6 (zinc finger protein 51)	NM_002125	2.96
major histocompatibility complex, class II, DR beta 5	NM_004741	2.96
nuclear and coiled-body phosphoprotein 1	NM_018128	2.96
hypothetical protein FLJ10534	NM_038250	2.95
Consensus includes gb:BF308250 /FEA=EST /DB_XREF=gi:11255432 /DB_XREF=est:601890308F1 /CLONE=IMAGE:4131673 /UG=Hs.315103 ESTs. Moderately similar to C	BF308250	2.95
proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki)	BC002684	2.94
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump), beta polypeptide, 56/58kD, isoform 2	NM_001693	2.94
manic fringe homolog (Drosophila)	AF36965	2.94
sperm associated antigen 9	NM_003971	2.93
apoptosis antagonizing transcription factor	AF063206	2.93
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 21kD	BC005876	2.93
HSPC003 protein	NM_014017	2.93
nuclear prelamin A recognition factor	NM_012336	2.93
ribosomal protein L5	NM_000969	2.92
sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase)	AV695711	2.92
soc-2 suppressor of clear homolog (C. elegans)	NM_007373	2.91
adenosine kinase	U90339	2.91
mitochondrial ribosomal protein S7	NM_015971	2.91
CD38 antigen (p45)	NM_001775	2.91

major histocompatibility complex, class I, C
 Consensus includes gb:AL353681 /DEF=Human DNA sequence from clone RP1-158P9 on chromosome 1. Contains a putative novel gene, a laminin receptor 1 (67kD, ribosom
 hypothetical protein FLJ10439
 adenosine deaminase
 membrane component, chromosome 11, surface marker 1
 chromosome 20 open reading frame 1
 arachidonate 5-lipoxygenase-activating protein
 fusion, derived from t(12;18) malignant liposarcoma
 casein kinase 2, alpha 1 polypeptide
 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 8 (19kD, PGIV)
 HLA-G histocompatibility antigen, class I, G
 potassium channel, subfamily K, member 12
 HSPC037 protein
 SH3 domain binding glutamic acid-rich protein like 3
 sphingosine-1-phosphatase
 vesicle-associated membrane protein 1 (synaptobrevin 1)
 ubiquitin-like 5
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily f, member 1
 thyroid receptor interacting protein 15
 Consensus includes gb:AK000185.1 /DEF=Homo sapiens cDNA FLJ20178 fls, clone COL09990. /FEA=mRNA /DB_XREF=gi:7020104 /UG=Hs.306389 Homo sapiens cDNA FI
 pericentriolar material 1
 CDC16 cell division cycle 16 homolog (S. cerevisiae)
 acute carrier family 1 (neutral amino acid transporter), member 5
 dynein, cytoplasmic, heavy polypeptide 1
 GDP-mannose pyrophosphorylase A
 ATX1 antioxidant protein 1 homolog (yeast)
 fragile X mental retardation, autosomal homolog 1
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily b, member 1
 H2A histone family, member X
 Consensus includes gb:AA46867 /FEA=EST /DB_XREF=gi:2216118 /DB_XREF=estab3803.s1 /CLONE=IMAGE:843101 /UG=Hs.284235 ESTs
 spectrin SH3 domain binding protein 1
 CCR4-NOT transcription complex, subunit 8
 proteasome (prosome, macropain) subunit, beta type, 4
 CGI-47 protein
 phosphoserine aminotransferase
 nuclear matrix protein NMP200 related to splicing factor PRP19
 transforming growth factor, beta 2
 replication factor C (activator 1) 5 (36.5kD)
 RNA binding motif protein 4
 splicing factor 3a, subunit 3, 60kD
 cytoskeleton-associated protein 1
 SH3-domain, GRB2-like, endophilin B2
 neural precursor cell expressed, developmentally down-regulated 5
 serine/threonine kinase 15
 DnaJ (Hsp40) homolog, subfamily C, member 8
 map kinase phosphatase-like protein MK-STYX
 CD789 antigen (immunoglobulin-associated beta)
 uridine monophosphate kinase
 hypothetical protein PRO1847
 chromosome 20 open reading frame 80
 interferon stimulated gene (20kD)
 similar to S. cerevisiae RER1
 adaptor-related protein complex 2, sigma 1 subunit
 eukaryotic translation initiation factor 4E-like 3
 eukaryotic translation initiation factor 5A
 Consensus includes gb:A591100 /FEA=EST /DB_XREF=gi:4600148 /DB_XREF=asttw01cd3.x1 /CLONE=IMAGE:2267044 /UG=Hs.113368 neuroendocrine secretory protein
 gb:M24669.1 /DEF=Human Ig rearranged H-chain V-region mRNA (C-D-JH6), complete cds. /FEA=mRNA /GEN=IGH@ /DB_XREF=gi:185200 /FL=gb:M24669.1
 Breakpoint cluster region protein, uterine leiomyoma, 1: barrier to autointegration factor
 cytoskeleton-associated protein 1
 RAD23 homolog A (S. cerevisiae)
 hypothetical protein FLJ20859
 membrane interacting protein of RGS16
 polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)

AK024838 2.90
 BC006351 2.90
 NM_000022 2.89
 NM_005698 2.89
 AF098158 2.89
 NM_001629 2.89
 S75762 2.89
 A1181318 2.89
 NM_014222 2.89
 M90666 2.88
 NM_022055 2.88
 BC003186 2.88
 NM_031286 2.88
 NM_030791 2.88
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 NM_024292 2.87
 AF231056 2.87
 NM_004236 2.87
 AK000185 2.86
 NM_006197 2.86
 NM_003903 2.86
 AF105230 2.86
 AB002323 2.86
 NM_013335 2.85
 NM_004045 2.85
 A1990766 2.85
 AK021419 2.85
 NM_002105 2.85
 AA48687 2.85
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 A1769416 2.84
 NM_002796 2.84
 NM_016014 2.84
 A189380 2.83
 NM_014502 2.83
 NM_003238 2.83
 NM_007370 2.83
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 NM_020145 2.81
 NM_004404 2.81
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 NM_014280 2.80
 NM_016086 2.80
 NM_000626 2.80
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 AF119855 2.79
 AL441988 2.79
 U88964 2.79
 NM_007033 2.78
 NM_004089 2.78
 AF047695 2.77
 NM_001870 2.76
 A1591100 2.76
 M24669 2.76
 AF044773 2.76
 BC005969 2.75
 BF572938 2.75
 NM_022734 2.75
 NM_016641 2.75
 AA679988 2.75

TNF-induced protein	BC005352	2.75
HLA-G histocompatibility antigen, class I, G	M90685	2.74
deoxyguanosine kinase	NM_001929	2.74
KIAA1228 protein	AJ708767	2.74
cut-like 1, CCAAT displacement protein (Drosophila)	NM_001913	2.74
endonuclease G	NM_004435	2.74
CREBBP/EP300 inhibitory protein 1	AF274951	2.74
damage-specific DNA binding protein 1 (127KD)	L40326	2.74
RAN binding protein 7	AL137335	2.73
sorting nexin 5	NM_014426	2.73
hydroxyacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit	U04627	2.73
transportin-SR	NM_012470	2.72
24-dehydrocholesterol reductase	NM_014762	2.72
microtubule-associated proteins 1A/1B light chain 3	AF183417	2.72
GDP dissociation inhibitor 2	NM_001494	2.72
bleomycin hydrolase	NM_000386	2.72
DNA segment, single copy probe LNS-CAILNS-CALI (deleted in polyposis)	A4814140	2.72
major histocompatibility complex, class II, DM alpha	X76775	2.72
adaptor-related protein complex 2, sigma 1 subunit	BC006337	2.72
UDP glycosyltransferase 2 family, polypeptide B15	NM_010176	2.72
Consensus includes gb:AW582267 /FEA=EST /DB_XREF=gi:7257316 /DB_XREF=gi:7257316 /DB_XREF=est:QVQV-ST0215-060100-083-c06 /UG=Hs.30695.1 Human DNA sequence from clone RP11	AW582267	2.72
MCHM4 minichromosome maintenance deficient 4 (S. cerevisiae)	A859865	2.71
laminin receptor 1 (87KD, ribosomal protein SA)	AL136306	2.71
KIAA0095 gene product	NM_014669	2.71
serine/threonine kinase 24 (STE20 homolog, yeast)	AA586774	2.71
RAB5A, member RAS oncogene family	NM_004162	2.71
solute carrier family 7 (cationic amino acid transporter, y+ system), member 5	A361227	2.70
src family associated phosphoprotein 2	AB018009	2.70
RD RNA-binding protein	AC003999	2.70
lymphocyte-specific protein tyrosine kinase	L03411	2.70
cytochrome c-1	U07236	2.70
core histone macroH2A2.2	NM_019116	2.70
deoxyhypusine synthase	NM_018649	2.70
hypothetical protein MGC10433	U26266	2.70
Consensus includes gb:BE988833 /FEA=EST /DB_XREF=gi:10579538 /DB_XREF=est:60164986 /F1 /CLONE=IMAGE:3933782 /UG=Hs.324648 Homo sapiens cDNA FLJ137	NM_024321	2.69
immunoglobulin lambda joining 3	BE988833	2.69
zinc finger protein 143 (clone pHZ-1)	X57812	2.69
3'-phosphoadenosine 5'-phosphosulfate synthase 1	AJ277546	2.69
Fanconi anemia, complementation group G	AF033026	2.69
RAB4, member RAS oncogene family	AC004472	2.69
postmeiotic segregation increased 2-like 3	NM_004578	2.69
gb:NM_016592.1 /DEF=Homo sapiens neuroendocrine secretory protein 55 (NESP55), mRNA, /FEA=mRNA /GEN=NESP55 /PROD=neuroendocrine secretory protein 55 /DB_XREF=gi:10579538 /DB_XREF=est:60164986 /F1 /CLONE=IMAGE:3933782 /UG=Hs.324648 Homo sapiens cDNA FLJ137	D38437	2.68
adenine phosphoribosyltransferase	NM_016592	2.68
hypothetical protein R32184_1	NM_000485	2.68
deoxyhypusine synthase	BC001648	2.68
BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)	NM_001930	2.68
voltage-dependent anion channel 1	AF081496	2.67
Rho GDP dissociation inhibitor (GDI) beta	AL151918	2.67
ribosomal protein L24	NM_001175	2.67
gb:NM_018604.1 /DEF=Homo sapiens hypothetical protein PRO1741 (PRO1741), mRNA, /FEA=mRNA /GEN=PRO1741 /PROD=hypothetical protein PRO1741 /DB_XREF=gi:10579538 /DB_XREF=est:60164986 /F1 /CLONE=IMAGE:1127826 /UG=Hs.132390 zinc finger protein 36 (KOX 18)	NM_000986	2.67
Consensus includes gb:AA653300 /FEA=EST /DB_XREF=gi:2569471 /DB_XREF=est:ag65c10.s1 /CLONE=IMAGE:1127826 /UG=Hs.132390 zinc finger protein 36 (KOX 18)	NM_018604	2.67
muscleblind-like (Drosophila)	AA653300	2.66
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1)	N31913	2.66
high-mobility group 20B	NM_002734	2.66
anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phospholipase A2)	AF288679	2.66
H2A histone family, member Y	BE869583	2.66
casein kinase 2, beta polypeptide	AF044286	2.66
zinc finger protein 43 (HITF8)	M30448	2.65
hypothetical protein MGC10715	NM_003423	2.65
mesenchymal stem cell protein DSC43	AL049650	2.65
proteasome (prosome, macropain) subunit, beta type, 9 (large multifunctional protease 2)	A1992095	2.65
adenylyate kinase 2	NM_002800	2.65
	W02312	2.65

mesenchymal stem cell protein DSC92	NM_016645	2.85
transcription factor CP2	U03494	2.85
polymerase (RNA) II (DNA directed) polypeptide 1 (14.5kD)	AL037557	2.84
polymerase (RNA) II (DNA directed) polypeptide D	NM_004805	2.84
RNA-binding protein (autoantigenic)	NM_016732	2.83
hepatocellular carcinoma-associated antigen 66	NM_016428	2.83
ADP-ribosylation factor 1	AA580004	2.83
v-ref-1 murine leukemia viral oncogene homolog 1	NM_002880	2.83
triosephosphate isomerase 1	NM_000385	2.83
seb4D	AL109955	2.83
polymerase (RNA) III (DNA directed) (62kD)	BC004424	2.83
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump), member D	AL568172	2.82
thioredoxin domain-containing	NM_030755	2.82
FK506 binding protein 1A (12kD)	NM_000801	2.81
cyclochrome b-245, alpha polypeptide	NM_000101	2.81
eukaryotic translation initiation factor 4E binding protein 1	AB046548	2.81
aldo-keto reductase family 1, member A1 (aldehyde reductase)	NM_008066	2.81
archelin 1	NM_001655	2.81
death associated protein 3	U16321	2.81
RuvB-like 1 (E. coli)	NM_003707	2.81
RNA binding motif protein, X chromosome	AA52524	2.80
fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome)	NM_023107	2.80
mutS homolog 5 (E. coli)	AF034759	2.80
dihydroipomide S-acyltransferase (E2 component of pyruvate dehydrogenase complex)	J03866	2.80
ribosomal protein L44	NM_021029	2.80
high-mobility group (nonhistone chromosomal) protein 14	NM_004965	2.80
Ras-GTPase activating protein SH3 domain-binding protein 2	NM_012297	2.59
60S ribosomal protein L30 isolog	NM_016304	2.59
AD021 protein	NM_016613	2.59
uncoupling protein 2 (mitochondrial, proton carrier)	U82819	2.59
60S acidic ribosomal protein PO	NM_016183	2.58
C-terminal binding protein 1	NM_001328	2.58
ATP synthase, H ⁺ transporting, mitochondrial F1 complex, gamma polypeptide 1	NM_005174	2.58
insulin induced gene 1	NM_005542	2.58
chloride intracellular channel 4	AF109196	2.58
ribosomal protein S25	AA888388	2.58
SEC13-like 1 (S. cerevisiae)	NM_030673	2.58
survival of motor neuron protein interacting protein 1	AB037701	2.58
CGI-150 protein	AF061730	2.57
hypothetical protein FLJ12287 similar to semaphorins	NM_022367	2.57
proline synthetase co-transcribed (bacterial homolog)	NM_007198	2.57
ribosomal protein S8	BC000524	2.57
breast carcinoma amplified sequence 4	NM_017843	2.57
phosphoglucanate dehydrogenase	NM_002631	2.57
transferrin receptor (p80, CD71)	NM_003234	2.57
neural precursor cell expressed, developmentally down-regulated 5	AI191427	2.57
enkephase promoting complex subunit 5	BF667271	2.57
MAP kinase-interacting serine/threonine kinase 1	BC002755	2.57
calnexin	L18887	2.56
gila maturation factor, gamma	NM_004877	2.56
ATPase, Na ⁺ /K ⁺ transporting, beta 3 polypeptide	U51478	2.56
sorting nexin 1	AF065484	2.55
Rho GDP dissociation inhibitor (GDI) alpha	BC005851	2.55
histidyl-RNA synthetase	NM_002109	2.55
platelet/endothelial cell adhesion molecule (CD31 antigen)	M37760	2.55
KIAA0123 protein	BF570122	2.55
ret finger protein	NM_018150	2.55
hypothetical protein FLJ10597	AF230394	2.55
interleukin 21 receptor	AF269133	2.55
ARF3 actin-related protein 3 homolog (yeast)	NM_005721	2.54
ADP-ribosylation factor 4	AL537042	2.54
heterogeneous nuclear ribonucleoprotein F	AI591354	2.54
phorbol-12-myristate-13-acetate-induced protein 1	AI857639	2.53

cAMP responsive element modulator	NM_001881	2.53
ubiquitin-conjugating enzyme E2 variant 1	U39361	2.53
immunoglobulin lambda locus	AF043586	2.53
chromosome condensation-related SMC-associated protein 1	AK022511	2.53
hypothetical protein MGC4368	NM_024510	2.53
fer (lipo/ies related) tyrosine kinase (phosphoprotein NCP94)	NM_005246	2.53
chromatin assembly factor 1, subunit A (p150)	NM_005483	2.52
estrogen-related receptor alpha	L39487	2.52
hypothetical protein MGC5242	NM_024033	2.52
glyceraldehyde-3-phosphate dehydrogenase	AK026525	2.52
hypothetical protein FLJ10702	NM_018184	2.51
valyl-tRNA synthetase 2	NM_006295	2.51
heat shock 90kD protein 1, alpha	AI862933	2.51
villin 2 (extrin)	BF663141	2.51
desitin (actin depolymerizing factor)	NM_021130	2.50
peptidylprolyl isomerase A (cyclophilin A)	NM_015640	2.50
PAL-1 mRNA-binding protein	NM_000284	2.50
pyruvate dehydrogenase (lipoamide) alpha 1	NM_016007	2.50
gb:NM_016007.1/DEF=Homo sapiens CGI-59 protein (LOC51625), mRNA, /FEA=mRNA /GEN=LOC51625 /PROD=CGI-59 protein /DB_XREF=gi:7706297 /UG=Hs.279887 C	NM_016007	2.50
protein phosphatase 1, regulatory (inhibitor) subunit 2	NM_006241	2.49
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 17 (72kD)	U59321	2.49
glycyl-tRNA synthetase	D30658	2.49
putative translation initiation factor	BF246436	2.49
putative translation initiation factor	Z97353	2.48
ribosomal protein L15	NM_024006	2.48
hypothetical protein IMAGE3455200	NM_017916	2.48
hypothetical protein FLJ20843	NM_024579	2.48
hypothetical protein FLJ23221	BC000924	2.48
NBR2	AL537707	2.48
putative translation initiation factor	U54829	2.47
regulator of G-protein signalling 16	NM_006503	2.47
proteasome (prosome, macropain) 26S subunit, ATPase, 4	AA167775	2.47
NRAS-related gene	NM_003340	2.47
ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast)	NM_030978	2.47
hypothetical protein similar to actin related protein 2/3 complex, subunit 5	AF189015	2.47
villin 2 (extrin)	NM_002949	2.46
mitochondrial ribosomal protein L12	BC004334	2.46
ribosomal protein S10	BC288967	2.46
RAD21 homolog (S. pombe)	NM_001428	2.46
endolase 1, (alpha)	NM_021633	2.46
kelch-like protein C3IP1	AL136932	2.45
inhibitor of growth family, member 1	AF044076	2.45
putative cyclin G1 interacting protein	NM_006349	2.45
CD184 antigen, salomuch	BF669455	2.45
ubiquinol-cytochrome c reductase hinge protein	NM_006004	2.45
aspartyl aminopeptidase	NM_012100	2.45
chaperonin containing TCP1, subunit 8 (theta)	NM_006585	2.44
superoxide dismutase 1, soluble (erythrocyte lateral sclerosis 1 (adult))	NM_000454	2.44
zinc finger protein	NM_015871	2.44
isocitrate dehydrogenase 3 (NAD+) beta	AF023266	2.44
sterol regulatory element binding transcription factor 2	NM_004599	2.43
phosphoribosyl pyrophosphate synthetase 1	NM_002764	2.43
KIAA0074 protein	D38553	2.43
hypothetical protein MGC4022	AI805286	2.43
uncharacterized bone marrow protein BM039	NM_018455	2.43
integral type I protein	BC000027	2.43
Consensus includes gb:M21388.1 /DEF=Human unproductively rearranged Ig mu-chain mRNA V-region (VD), 5' end, clone mu-3A1A, /FEA=mRNA /GEN=IGHM /DB_XREF=g	M21388	2.43
desmoglein 1	NM_001942	2.43
BCL2-associated athanogene	AF116273	2.43
general transcription factor II, i	NM_001518	2.43
ubiquinol-cytochrome c reductase, Rieske iron-sulfur polypeptide 1	BC000649	2.42
histone H2A.F/Z variant	BF343852	2.42
ribosomal protein S2	AA630314	2.42

Consensus includes gb:BE963164 /FEA=EST /DB_XREF=gi:11766582 /DB_XREF=est:601656973R1 /CLONE=IMAGE:3865650 /UG=Hs.2186 eukaryotic translation elongation. BE963164 2.42

IGF-II mRNA-binding protein 3 AU160004 2.42

Consensus includes gb:AL121975 /DEF=human DNA sequence from clone RP3-42ZB11 on chromosome 6p11.2-12.3 Contains ESTs, STSs and GSSs. Contains part of the P AL121975 2.42

proteolipid protein 2 (colonic epithelium-enriched) BC005882 2.42

peptidylprolyl isomerase A (cyclophilin A) NM_030666 2.41

serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1 NM_014402 2.41

low molecular mass ubiquitinone-binding protein (9.5kD) NM_016576 2.41

GNP2 for guanosine monophosphate reductase isolog NM_007096 2.41

clathrin, light polypeptide (Lca) U38664 2.40

postmeiotic segregation increased 2-like 8 AV007161 2.40

SRY (sex determining region Y)-box 2 N29889 2.40

UBX domain-containing 2 A1628925 2.40

KIAA1899 protein NM_020381 2.40

candidate tumor suppressor protein NM_014183 2.40

HSPC162 protein AA502912 2.39

KIAA0906 protein NM_004526 2.39

MCM2 minichromosome maintenance deficient 2, mitotin (S. cerevisiae) BC003574 2.39

T-cell leukemia/lymphoma 1A NM_002573 2.39

platelet-activating factor acetylhydrolase, isoform 1b, gamma subunit (29kD) NM_005759 2.39

proteasome (prosome, macropain) activator subunit 3 (PA28 gamma: Ki) AL518328 2.39

basic leucine-zipper protein BZAP45 NM_020187 2.39

DC12 protein BF342707 2.38

pre-mRNA processing factor 31 homolog (yeast) AF074723 2.38

RNA polymerase II transcriptional regulation mediator (Med6, S. cerevisiae, homolog of) NM_003712 2.38

protein phosphatase 1, regulatory subunit 7 NM_003730 2.38

ribonuclease 6 precursor NM_003932 2.38

suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein) NM_003932 2.38

zinc finger protein 195 AA476267 2.38

kinasin 2 (60-70kD) AF164622 2.38

golgin-47 NM_001786 2.37

cell division cycle 2, G1 to S and G2 to M BG034328 2.37

transcription factor Op-2 (E2F dimerization partner 2) AF683882 2.37

Rho-associated, coiled-coil containing protein kinase 1 gb:AF274954.1 /DEF=Homo sapiens PNAS-29 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-29 /DB_XREF=gi:12751070 /UG=Hs.326791 Homo sapiens PNAS-29 mRNA /AF274954 2.37

HSPC002 protein NM_015382 2.37

gb:NM_004682.1 /DEF=Homo sapiens PC4 and SFRS1 interacting protein 2 (PSIP2), mRNA. /FEA=mRNA /GEN=PSIP2 /PROD=PC4 and SFRS1 interacting protein 2 /DB_XF NM_004682 2.37

hypothetical protein MGC2722 BF975273 2.36

uncharacterized hypothalamus protein HT009 NM_018470 2.36

lipok acid synthetase BF056778 2.36

dUTP pyrophosphatase U90223 2.36

mitochondrial ribosomal protein L17 AK026657 2.36

ribosomal protein S15a NM_001019 2.36

dynamitin 1 (p150, glued homolog, Drosophila) NM_004082 2.35

polymerase (RNA) II (DNA directed) polypeptide G NM_002696 2.35

H2B histone family, member A BC000893 2.35

major histocompatibility complex, class I, A A4573862 2.35

activity-dependent neuroprotector NM_015339 2.35

mitochondrial ribosomal protein S17 NM_015969 2.35

Consensus includes gb:BE250348 /FEA=EST /DB_XREF=gi:3120458 /DB_XREF=est:600943342T1 /CLONE=IMAGE:2959811 /UG=Hs.326249 Homo sapiens mRNA; cDNA [BE250348 2.35

FXYD domain-containing ion transport regulator 5 NM_014164 2.34

actin related protein 2/3 complex, subunit 5 (16 kD) AL516350 2.34

matrin 3 A4577695 2.34

general transcription factor II, i BC004472 2.34

Wolf-Hirschhorn syndrome candidate 1 BE793789 2.34

casein kinase 2, alpha prime polypeptide NM_001896 2.34

nuclear receptor co-repressor 1 AA085748 2.34

synovial sarcoma translocation, chromosome 18 NM_005637 2.33

eukaryotic translation elongation factor 1 alpha 1 AL515273 2.33

casein kinase 1, alpha 1 NM_001892 2.33

methylcrotonoyl-Coenzyme A carboxylase 2 (beta) AB050049 2.33

prostaglandin D2 synthase, hemopoietic NM_014485 2.33

stem-loop (histone) binding protein NM_006527 2.33

isocitrate dehydrogenase 3 (NAD+) alpha A1826060 2.33

hypothetical protein FLJ21617	NM_030897	2.33
ribosomal protein S12	AI799007	2.33
kinesin-like 6 (mitotic centromere-associated kinesin)	U63743	2.33
chromosome condensation 1	X06130	2.33
ribophorn II	BC003560	2.33
protein phosphatase 6, catalytic subunit	NM_002721	2.33
splicing factor, arginine/serine-rich 7 (55kD)	AA524053	2.33
Wolf-Hirschhorn syndrome candidate 1	AF083389	2.33
serum/glucocorticoid regulated kinase-like	NM_013257	2.33
guanine nucleotide binding protein (G protein), alpha 13	NM_006572	2.32
MYG1 protein	NM_021640	2.32
FtsJ homolog 2 (E. coli)	AK024635	2.32
accessory proteins BAP31/BAP29	NM_005745	2.32
TRAF family member-associated NFKB activator	NM_004180	2.32
cytochrome b5 outer mitochondrial membrane precursor	NM_030579	2.32
hypothetical protein MGC4504	NM_024111	2.32
pyridoxal (pyridoxine, vitamin B6) kinase	NM_003681	2.32
protease, serine, 22	NM_022119	2.31
ribosomal protein L4	A053886	2.31
zinc finger protein 225	AC074331	2.31
general transcription factor IIC, polypeptide 3 (102kD)	NM_012086	2.31
SWI3 suppressor of mit two 3 homolog 1 (yeast)	BG338532	2.31
hypothetical protein FLJ20436	AK022732	2.31
mitogen-activated protein kinase kinase 7	AF218074	2.31
D-dopa-chrome tautomerase	NM_001355	2.31
translocase of outer mitochondrial membrane 20 (yeast) homolog	NM_014755	2.31
KIAA0255 gene product	A141892	2.31
RAD23 homolog B (S. cerevisiae)	AL527385	2.31
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	NM_000351	2.31
HSPC053 protein	NM_014151	2.31
6-phosphogluconolactonase	NM_012088	2.31
mannosidase, alpha, class 1A, member 1	NM_005907	2.31
proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; Ki)	AA758755	2.31
adenylosuccinate lyase	NM_000026	2.30
alkylglycerone phosphate synthase	NM_003659	2.30
dendritic cell protein	BC001465	2.30
HBS1-like (S. cerevisiae)	NM_006360	2.30
hypothetical protein FLJ12806	NM_022831	2.30
amyloid beta (A4) precursor-like protein 2	BG427393	2.29
X-prolyl aminopeptidase (aminopeptidase P)-like	W07689	2.29
PK domain containing 16	NM_006523	2.29
ring finger protein 10	NM_022114	2.29
high-mobility group (nonhistone chromosomal) protein 1	NM_014868	2.29
WW domain binding protein 11	BE311760	2.29
diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)	AF118023	2.29
ATP-dependent interferon responsive	NM_001845	2.29
neutrophil cytosolic factor 1 (47kD, chronic granulomatous disease, autosomal 1)	NM_022371	2.28
peptidyl prolyl isomerase H (cyclophilin H)	AW072386	2.28
tryptophanyl-tRNA synthetase	NM_006347	2.28
major histocompatibility complex, class I, C	NM_004184	2.28
KIAA1228 protein	BC004489	2.28
postmeiotic segregation increased 2-like 1	BC001224	2.28
eukaryotic translation initiation factor 28, subunit 2 (beta, 39kD)	D38435	2.28
DKFZP586A011 protein	NM_014239	2.28
H3 histone family, member L	NM_015416	2.28
Finkel-Biskamp-Reilly murine sarcoma virus (FBR-MuSV) ubiquitously expressed (fox derived); ribosomal protein S30	NM_003537	2.27
hypothetical protein PRO1635	NM_001987	2.27
hypothetical protein MGC2718	NM_016589	2.27
seven in absentia homolog 2 (Drosophila)	NM_024067	2.27
protein tyrosine phosphatase, non-receptor type 22 (lymphoid)	U76248	2.27
nuclear transcription factor Y, gamma	NM_015967	2.27
hypothetical protein FLJ23209	U82296	2.27
	NM_024895	2.27

hypothetical protein FLJ12549	NM_024844	2.27
BRCA1 associated protein	AF035620	2.27
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72kD)	AW188131	2.26
glyoxylate reductase/hydroxypyruvate reductase	NM_012203	2.26
collagen, type VI, alpha 1	M20776	2.26
protein kinase domains containing protein similar to phosphoprotein C8FW	NM_021158	2.26
progesterin induced protein	AF006010	2.25
HSPC160 protein	NM_014182	2.25
zinc finger protein 9 (a cellular retroviral nucleic acid binding protein)	NM_003418	2.25
H2A histone family, member Y	AF044286	2.25
polymerase (DNA-directed), delta 4	NM_021173	2.25
divalent cation tolerant protein CUTA	AF230924	2.25
hypothetical protein AF140225	NM_030799	2.25
immunoglobulin lambda locus	AF047245	2.25
isocitrate dehydrogenase 3 (NAD+) beta	AF023265	2.25
gb:D13287.1/DEF=Human mRNA for alternative splicing product of glucocerebrosidase pseudogene, complete cds. /FEA=mRNA (PROD=glucocerebrosidase /DB_XREF=gi:4	AB049944	2.25
mitochondrial ribosomal protein S11	BF976260	2.24
transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)	A1701949	2.24
heterogeneous nuclear ribonucleoprotein K	AA684258	2.24
heterogeneous nuclear ribonucleoprotein C (C1/C2)	U93240	2.24
karyopherin alpha 4 (importin alpha 3)	NM_014303	2.24
pescedillo homolog 1, containing BRCT domain (zebrafish)	NM_003315	2.24
DnaJ (Hsp40) homolog, subfamily C, member 7	BC003584	2.24
dihydrofolate reductase	U49188	2.24
tumor differentially expressed 1	NM_021968	2.24
H4 histone family, member E	NM_001628	2.24
aldo-keto reductase family 1, member B1 (aldose reductase)	BC006301	2.24
anaphase promoting complex subunit 5	BF689284	2.24
proliferation-associated ZC4, 38kD	NM_017654	2.24
hypothetical protein FLJ20512	BC000425	2.24
protein disulfide isomerase related protein (calcium-binding protein, intestinal-related)	AF263279	2.23
CD164 antigen, sialomucin	NM_005863	2.23
neuroepithelial cell transforming gene 1	BC000436	2.23
endosulfine alpha	NM_006414	2.23
ribonuclease P (38kD)	NM_012394	2.23
prefoldin 2	NM_012394	2.23
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capping protein (actin filament) muscle Z-line, beta	NM_007317	2.23
ribosomal protein S10	NM_004930	2.23
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 31kD	NM_001014	2.23
KIAA0310 gene product	BC004443	2.23
ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)-like 1	BC001404	2.23
clathrin and palatol associated transmembrane protein 1	AL359763	2.23
3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (soluble)	BC004865	2.23
stathin, RNA binding protein (Drosophila)	NM_002130	2.23
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 16kD	AL136801	2.23
bone morphogenetic protein 7 (osteogenic protein 1)	NM_001694	2.22
Consensus includes gb:AW054828 /FEA=EST /DB_XREF=gi:5920529 /DB_XREF=est:ws60c12.x1 /CLONE=IMAGE:2501590 /UG=Hs.58220 Homo sapiens cDNA: FLJ23005	M60316	2.22
proteasome (prosome, macropain) 26S subunit, ATPase, 1	NM_002802	2.22
translation initiation factor IF2	NM_015904	2.22
phosphomannomutase 1	NM_002676	2.22
mitogen-activated protein kinase kinase 7	AB009358	2.22
hypothetical protein MGC2594	NM_024050	2.22
small inducible cytokine subfamily B (Cys-X-Cys), member 11	AF030514	2.21
Rho GDP dissociation inhibitor (GDI) alpha	D13989	2.21
hypothetical protein FLJ11301	AL136897	2.21
serologically defined colon cancer antigen 16	BC001149	2.21
PEF protein with a long N-terminal hydrophobic domain (pelfin)	NM_012392	2.21
ribonucleotide reductase M2 polypeptide	BC001886	2.21
hypothetical protein FLJ21617	AK025270	2.21
Rho GDP dissociation inhibitor (GDI) alpha	NM_004309	2.21
CCR4-NOT transcription complex, subunit 8	AF180476	2.21

intercellular adhesion molecule 3
 KIAA0092 gene product
 HSPC144 protein
 tubulin, alpha 1 (testis specific)
 postmeiotic segregation increased 2-like 8
 mitochondrial ribosomal protein S18B
 5-enolmaltase-4-carboxamide ribonucleotide formyltransferase/IMP cyclohydrolase
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 ribosomal protein S21
 poly(rC) binding protein 2
 NOL1R protein
 neutrophil cytosolic factor 1 (47kD, chronic granulomatous disease, autosomal 1)
 programmed cell death 8 (apoptosis-inducing factor)
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 methionine adenosyltransferase II, alpha
 TIA1 cytosolic granule-associated RNA binding protein
 apoptosis related protein APR-3
 ADP-ribosyltransferase (NAD+; poly (ADP-ribose) polymerase)
 FUS interacting protein (serine-arginine rich) 2
 ubiquitination factor E48 (UFD2 homolog, yeast)
 CDC37 cell division cycle 37 homolog (S. cerevisiae)
 hypothetical protein FLJ11200
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 polymerase (RNA) III (DNA directed) (62kD)
 nuxix (nucleoside diphosphate linked moiety X)-type motif 1
 hypothetical protein DKFp761H221
 ATPase, H+ transporting, lysosomal (vacuolar proton pump), member J
 kinesin-like 6 (mitotic centromere-associated kinesin)
 gb-NM_016402.1 /DEF=Homo sapiens hypothetical protein (HSPC140), mRNA. /FEA=mRNA (GEN=HSPC140 /PROD=hypothetical protein /DB_XREF=gi:7705472 /UG=Hs.27
 hypothetical protein FLJ10479
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 35
 hypothetical protein FLJ20257
 KIAA1513 protein
 CDC8 cell division cycle 8 homolog (S. cerevisiae)
 H2A histone family, member C
 H3 histone family, member B
 BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)
 inositol polyphosphate-5-phosphatase, TSKD
 thousand and one amino acid protein kinase
 ribosomal protein S3A
 hypothetical protein FLJ11301
 CASP2 and RIPK1 domain containing adaptor with death domain
 karyopherin (importin) beta 1
 protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform
 FK506 binding protein 2 (13kD)
 voltage-dependent anion channel 3
 hypothetical gene CG018
 glutamyl-prolyl-4-aminopeptidase
 similar to yeast BET3 (S. cerevisiae)
 ferritin, heavy polypeptide pseudogene 1
 homolog of yeast MCM10; hypothetical protein PRO2249
 KIAA1049 protein
 ribosomal protein L35a
 hypothetical protein FLJ21610
 hypothetical protein FLJ20312
 synrophin, beta 1 (dystrrophin-associated protein A1, 59kD, basic component 1)
 mitochondrial ribosomal protein L32
 UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 3
 chromosome 11 open reading frame2
 nucleolar and coiled-body phosphoprotein 1
 hypothetical protein
 hypothetical protein FLJ12298
 3-hydroxy-3-methylglutaryl-Coenzyme A reductase

gb:NM_017445.1 /DEF=Homo sapiens H2B histone family, member S (H2BFS), mRNA. /FEA=CDS /GEN=H2BFS /PROD=H2B histone family, member S /DB_XREF=gi:11036	NM_017445
dyskeratosis congenita 1, dyskerin	NM_001363
NAD kinase	BE551347
chondroitin sulfate proteoglycan 6 (barnacin)	BF795297
calcium-regulated heat-stable protein (24kD)	NM_014316
goli transport complex 1 (90 kDa subunit)	NM_006348
KIAA1858 protein	AL060102
potassium channel, subfamily K, member 1 (TWIK-1)	U90065
solute carrier family 8 (neurotransmitter transporter, GABA), member 13	NM_016615
hypothetical protein	AX000828
postmitotic segregation increased 2-like 8	AA243508
actin, gamma 1	NM_005394
lymphocyte adaptor protein	AW190090
carboxylate (N-acetylglucosamine 6-O) sulfotransferase 6	NM_005475
jumping translocation breakpoint	NM_021615
cytochrome b5 outer mitochondrial membrane precursor	BC004239
ATP-binding cassette, sub-family F (GCN20), member 1	AW235051
SVC (Src homology 2 domain-containing) transforming protein 1	NM_001090
RAD51 homolog C (S. cerevisiae)	A091079
hypothetical protein	AF029669
Consensus includes gb:S74774.1 /DEF=p59Yn(T)=OKT3-induced calcium influx regulator human, Jurkat J6 T cell line. /FEA=CDS /PROD=lysozyme kin. S74774	BE732345
hematopoietic cell-specific Lyn substrate 1	NM_005335
RAP1, GTP-GDP dissociation stimulator 1	X53465
pne-B-cell colony-enhancing factor	BF575514
2,5'-oligoadenylate synthetase 1 (40-46 kD)	NM_002534
hypothetical protein FLJ10803	A1765280
adaptor protein with pleckstrin homology and src homology 2 domains	NM_020979
hypothetical protein	NM_018986
prostate tumor over expressed gene 1	AL046054
chromodomain protein, Y chromosome-like	NM_004824
postmitotic segregation increased 2-like 9	NM_005395
mitotic control protein dis3 homolog	NM_014953
guanine nucleotide binding protein (G protein), beta 5	NM_006578
FK506 binding protein 1A (12kD)	A1936789
eukaryotic transition initiation factor 3, subunit 5 (epsilon, 47kD)	NM_003754
cystathyl-RNA synthetase	A1769685
pyruvate-5-carboxylate synthetase (glutamate gamma-semialdehyde synthetase)	NM_002860
survival of motor neuron protein interacting protein 1	A8037703
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4 (9kD, MLRQ)	NM_002489
HSPC023 protein	NM_014047
H2B histone family, member R	NM_021058
PI-3-kinase-related kinase SMG-1	BG256504
myosin light chain 1 slow s	NM_002475
RNA binding motif protein 5A	NM_005105
immediate early protein	NM_004907
nuclear cap binding protein subunit 2, 20kD	NM_007382
phosphatidylinositol glycan, class F	NM_002843
BCL2-associated antagonist 5	NM_004873
chromosome 2 open reading frame 3	AC005034
KIAA1393 protein	AB037814
prostatic binding protein	BE969671
H2B histone family, member K	NM_003525
acid phosphatase 1, soluble	NM_004300
Consensus includes gb:BF796470 /FEA=EST /DB_XREF=gi:12101524 /DB_XREF=est:60225959F1 /CLONE=IMAGE:4342999 /UG=Hs.284275 Homo sapiens PAX2 mRNA.	BF796470
FK506 binding protein 5	NM_004117
hypothetical protein FLJ20989	NM_023080
similar to S. pombe dim1+	NM_008701
eNOS interacting protein	NM_015953
ribosomal protein S4, X-linked	NM_001007
CD24 antigen (small cell lung carcinoma cluster 4 antigen)	L33930
Wilms' tumour 1-associating protein	BC000383
KIAA0625 protein	N64643

synaptic nuclei expressed gene 2	NM_015180	2.03
stearoyl-CoA desaturase (delta-9-desaturase)	BC005807	2.03
peptide deformylase-like protein	NM_022341	2.03
WD repeat domain 6	NM_018031	2.03
atrophin-1 interacting protein 1; actin receptor interacting protein 1	AC004990	2.03
catalase	AY028632	2.03
ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R)	NM_001419	2.03
SWAP-70 protein	AB014540	2.03
serologically defined colon cancer antigen 16	BC001148	2.03
transformation/transcription domain-associated protein	AC004893	2.03
tumor necrosis factor receptor superfamily, member 17	NM_001192	2.03
LIM and SH3 protein 1	NM_006148	2.03
nuclear cap binding protein subunit 1, 80kD	BC001450	2.03
KIAA1014 protein	AB023231	2.03
Wolff-Hirschhorn syndrome candidate 1	BF111870	2.03
DnaJ (Hsp40) homolog, subfamily A, member 2	AF011793	2.03
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit g	AL050277	2.03
omithine decarboxylase 1	NM_002539	2.03
defender against cell death 1	NM_001344	2.02
H2A histone family, member Z	NM_002106	2.02
O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase)	U77413	2.02
protease, serine, 25	AF184911	2.02
O-6-methylguanine-DNA methyltransferase	NM_002412	2.02
keratin, hair, acidic.3A	NM_004138	2.02
capillary renal cell carcinoma (translocation-associated)	BC004913	2.02
guanine nucleotide binding protein (G protein), beta 5	AF300650	2.02
calumenin	AF257659	2.02
GCM1 general control of amino-acid synthesis 1-like 1 (yeast)	D86973	2.02
cyclin D3	NM_001760	2.02
actin, gamma 1	AL515810	2.02
ribosomal protein L7a	NM_000972	2.02
B-cell translocation gene 1, anti-proliferative	NM_001731	2.01
KIAA1608 protein	NM_024820	2.01
stearoyl-CoA desaturase (delta-9-desaturase)	AA678241	2.01
Consensus includes gb:AL137162 /DEF=Human DNA sequence from clone RP5-843L14 on chromosome 20. Contains ESTs, STSs and GSSs. Contains a novel gene and the vitamin A responsive; cytoskeleton related	AL137162	2.01
KIAA0225 protein	N92494	2.01
chromosome 11 open reading frame 10	AW206115	2.01
eukaryotic translation elongation factor 2	NM_014206	2.01
polymerase (RNA) II (DNA directed) polypeptide C (33kD)	A004246	2.01
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 9 (39kD)	AC004382	2.01
proteasome (prosome, macropain) activator subunit 2 (PA28 beta)	AF050641	2.01
fibroblast growth factor (acdc) intracellular binding protein	NM_002818	2.01
hippocalcin-like 1	NM_004214	2.01
ras homolog gene family, member H	NM_002149	2.00
non-canonical ubiquitin conjugating enzyme 1	NM_004310	2.00
similar to hypothetical protein MNCb-2386	NM_016021	2.00
hypothetical protein FLJ20546	AA421957	2.00
metallothionein 2A	NM_017872	2.00
coated vesicle membrane protein	NM_005953	2.00
NOL1R protein	NM_018044	2.00
malic enzyme 2, NAD(+)-dependent, mitochondrial	BC000147	2.00
PRP4/STK/WD splicing factor	U82756	2.00
putative dimethyladenosine transferase	W87688	2.00
tumor necrosis factor receptor superfamily, member 6	X63493	2.00
mitogen-activated protein kinase kinase 3	AA1780381	2.00
delta sleep inducing peptide, immunoreactor	NM_004089	2.00
hypothetical protein	BE314601	2.00
survival of motor neuron protein interacting protein 1	AB037702	1.99
ubiquitin A-52 residue ribosomal protein fusion product 1	NM_015956	1.99
olfactory receptor, family 1, subfamily K, member 1	AF348700	1.99
forkhead box M1	NM_018635	1.99
	NM_021953	1.99

anhydrin 1, erythrocytic	AF005213	1.99
RAE1 RNA export 1 homolog (S. pombe)	U65943	1.99
hypothetical protein FLJ20244	NM_017722	1.99
NHP2 non-histone chromosome protein 2-like 1 (S. cerevisiae)	NM_005008	1.98
nudix (nucleoside diphosphate linked moiety X)-type motif 4	NM_019094	1.98
heterogeneous nuclear ribonucleoprotein C (C1/C2)	NM_004500	1.98
H2A histone family, member Y	NM_004893	1.98
sperm associated antigen 9	AB011088	1.98
myeloid cell leukemia sequence 1 (BCL2-related)	BF594446	1.98
CDC8 cell division cycle 6 homolog (S. cerevisiae)	U77949	1.98
cell division cycle 2, G1 to S and G2 to M	D88357	1.98
calcium/calmodulin-dependent protein kinase IG	AW016039	1.97
Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse)	AJ276888	1.97
BCL2-associated athanogene	NM_004323	1.97
ubiquinol-cytochrome c reductase core protein II	AV72738.1	1.97
Link guanine nucleotide exchange factor II	NM_016339	1.97
core-binding factor, beta subunit	AF284326	1.97
DAZ associated protein 2	AL534321	1.97
hypothetical protein FLJ22382	NM_025045	1.97
general transcription factor IIH, polypeptide 4 (52KD subunit)	NM_001517	1.97
tyrosine kinase 2	NM_003331	1.97
phosphoprotein enriched in astrocytes 15	BC002426	1.97
p47	NM_016143	1.97
hypothetical protein MGC2655	NM_024339	1.97
protein-L-isopartate (D-aspartate) O-methyltransferase	M83008	1.97
secretory carrier membrane protein 3	NM_005698	1.97
hypothetical nuclear factor SBB122	NM_020395	1.96
DnaJ (Hsp40) homolog, subfamily B, member 6	AF080569	1.96
UDP-GlcNAc:beta-Gal beta-1,3-N-acetylglucosaminyltransferase 1	NM_006577	1.96
protein kinase C and casein kinase substrate in neurons 2	NM_007229	1.96
H2A histone family, member A	NM_021062	1.96
hypothetical protein FLJ12860	NM_025152	1.96
DnaJ (Hsp40) homolog, subfamily C, member 8	AA843895	1.96
gb.BC000285.1 /DEF-Homo sapiens, clone MGC:3182, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:3182) /DB_XREF=gi:12653008 /UG=Hs.31541	W74577	1.96
ubiquitin-conjugating enzyme HBUCE1	M15887	1.96
diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)	NM_007282	1.96
RNA-binding protein regulatory subunit	AA35828	1.96
samlocalin 2	NM_002834	1.96
proteasome (prosome, macropain) subunit, beta type, 10	NM_002801	1.96
hypothetical protein FLJ20552	NM_017876	1.96
KIAA1033 protein	AK001657	1.96
ATP-binding cassette, sub-family E (ABCF), member 1	A002002	1.96
protein transport protein SEC81 alpha subunit isoform 1	NM_013336	1.96
estrogen-related receptor alpha	NM_004451	1.95
PTD018 protein	NM_016125	1.95
KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 1	NM_008801	1.95
pyruvate dehydrogenase (liponamide) beta	M34055	1.95
hypothetical protein MGC2650	NM_024108	1.95
hypothetical protein FLJ20707	NM_017938	1.95
signal sequence receptor, alpha (translocin-associated protein alpha)	A016620	1.95
solute carrier family 18 (monocarboxylic acid transporters), member 1	NM_003051	1.95
myosin, light polypeptide, regulatory, non-sarcomeric (20KD)	NM_006471	1.95
asparaginylglucosaminidase	W27131	1.95
hypothetical protein FLJ10326	NM_018060	1.95
protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), alpha isoform	NM_002717	1.95
ATPase, Ca++ transporting, plasma membrane 1	M65541	1.95
Sec23 homolog A (S. cerevisiae)	NM_006364	1.95
eukaryotic translation elongation factor 1 beta 2	NM_001959	1.94
replication factor C (activator 1) 4 (37KD)	NM_002916	1.94
proteasome (prosome, macropain) subunit, alpha type, 5	NM_002780	1.94
peroxiredoxin 2	NM_005809	1.94
NOL1R protein	A1284845	1.94

hypy-IRNA synthetase	AF285758	1.94
glutaredoxin (thioltransferase)	AF162789	1.94
aldolase A, fructose-bisphosphate	AK026577	1.94
JAK binding protein	AB005043	1.94
hypothetical protein FLJ11856	AK021918	1.94
glutathione S-transferase M1	NM_000561	1.94
KIAA0211 gene product	NM_014630	1.94
X-ray repair complementing defective repair in Chinese hamster cells 5 (double-strand-break rejoining; Ku autoantigen, 80kD)	J04977	1.94
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5 (13kD, B13)	NM_005000	1.93
diaphanous homolog 1 (Drosophila)	AF051782	1.93
hypothetical protein FLJ20517	NM_017859	1.93
galactosidase, beta 1	NM_000404	1.93
mannose phosphate isomerase	NM_002435	1.93
hypothetical protein FLJ10374	NM_018074	1.93
glutathione peroxidase 4 (phospholipid hydroperoxidase)	NM_002085	1.93
phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole synthetase	NM_000819	1.93
arsenate resistance protein ARS2	BE646076	1.93
Rho-associated, coiled-coil containing protein kinase 2	D87831	1.93
H2B histone family, member H	NM_003523	1.93
cell division cycle 27	NM_001256	1.93
vesicle docking protein p115	NM_003715	1.93
KIAA0174 gene product	NM_014761	1.92
KIAA0603 gene product	NM_014832	1.92
Consensus includes gb:AV715578 /FEA=EST /DB_XREF=est:AV715578 /CLONE=DCBBJ/G09 /UG=Hs.21056 Homo sapiens cDNA: FLJ21366 fls. cdc AV715578	NM_014832	1.92
Consensus includes gb:AL121916 /DEF=Human DNA sequence from clone RP11-189G13 on chromosome 20. Contains an RPL7A (60S ribosomal protein L7A) (SURF3) pseud	NM_014832	1.92
chromosome 1 open reading frame 16	NM_014837	1.92
cystathionine-beta-synthase	BE613178	1.92
hypothetical protein	NM_016618	1.92
sin3-associated polypeptide, 30kD	BF247098	1.92
polymerase I and transcript release factor	AF312393	1.92
RBP1-like protein	AA887480	1.92
signal peptidase 12kDa	NM_014041	1.92
aldolase A, fructose-bisphosphate	NM_000034	1.92
Ops-interacting protein 2	AL050353	1.92
stau6n, RNA binding protein (Drosophila)	NM_004602	1.92
glutathione synthetase	NM_000178	1.92
interferon regulatory factor 7	NM_004030	1.92
N-myc downstream-regulated gene 3	BE300252	1.92
heterogeneous nuclear ribonucleoprotein A3	AA528233	1.92
heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)	BC003621	1.92
cyclin-dependent kinase 5	NM_004935	1.91
CD47 antigen (Rn-related antigen, integrin-associated signal transducer)	Z25521	1.91
microtubule-associated protein, RP/EB family, member 1	NM_004935	1.91
gb:AF660511.1 /DEF=Homo sapiens clone 016b10 My016 protein mRNA, complete cds. /FEA=mrna /PROD=My016 protein /DB_XREF=gi:12001971 /UG=Hs.181634 Homo	NM_012325	1.91
SET translocation (myeloid leukemia-associated)	M93851	1.91
ATPase, H+ transporting, lysosomal (vacuolar proton pump) membrane sector associated protein M8-9	NM_005765	1.91
PWP2 periodic tryptophan protein homolog (yeast)	U56085	1.91
ubiquitin carboxyl-terminal esterase L1 (ubiquitin thioesterase)	NM_004181	1.91
POU domain, class 2, associating factor 1	NM_008235	1.91
McKusick-Kaufman syndrome	NM_018848	1.91
DKFZp568D133 protein	AL050050	1.91
guanine monophosphate synthetase	NM_003875	1.91
cysteinyl-IRNA synthetase	NM_001751	1.91
DEAD/H (Asp-Glu-Ala-AspHis) box polypeptide 30	NM_014966	1.91
synplektin; Huntingtin interacting protein 1	Y10931	1.91
I-complex 1	BC000665	1.91
opamine receptor D2	M30625	1.91
hypothetical protein FLJ20764	NM_017955	1.91
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 16kD	M62762	1.91
gamma-tubulin complex protein 2	NM_006659	1.91
Consensus includes gb:AI653730 /FEA=EST /DB_XREF=gi:4737709 /DB_XREF=est:w63612.x1 /CLONE=IMAGE:2307791 /UG=Hs.279009 matrix Gla protein	AI653730	1.91
CHK2 checkpoint homolog (S. pombe)	BC004207	1.91
transcription factor-like 4	BF056105	1.90

electron-transfer-flavoprotein, beta polypeptide
 Sjogren's syndrome nuclear autoantigen 1
 Integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)
 agrin
 proteasome (prosome, macropain) 26S subunit, non-ATPase, 13
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1
 calcium channel, voltage-dependent, beta 1 subunit
 RAD17 homolog (S. pombe)
 SMT3 suppressor of mit two 3 homolog 1 (yeast)
 EBNA-2 co-activator (100kD)
 ring finger protein 1
 signal recognition particle 72kD
 phosphatidylserine synthase 1
 c-Mpl binding protein
 Consensus includes gb:H97931 /FEA=EST /DB_XREF=gi:118816 /DB_XREF=est:yx04c02.s1 /CLONE=IMAGE:260738 /UG=Hs.173108 Homo sapiens cDNA: FLJ21897 fls, t:H97931
 Consensus includes gb:H97931 /FEA=EST /DB_XREF=gi:118816 /DB_XREF=est:yx04c02.s1 /CLONE=IMAGE:260738 /UG=Hs.173108 Homo sapiens cDNA: FLJ21897 fls, t:H97931
 interferon-related developmental regulator 1
 phosphodiesterase 8D, cGMP-specific, rod, delta
 sorting nexin 3
 similar to S. cerevisiae SSM4
 CTP synthase
 neuronal protein 17.3
 erythrocyte membrane protein band 7.2 (stomatilin)
 KIAA0152 gene product
 ubiquitin carboxyl-terminal esterase L3 (ubiquitin thiolesterase)
 KIAA0999 protein
 GTPase GES; REM protein
 heat shock 70kD protein 9B (mortalin-2)
 CD53 antigen
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)
 microtubule-associated protein like echinoderm EMAP
 hypothetical protein FLJ13449
 adenylate kinase 2
 eukaryotic translation initiation factor 2B, subunit 5 (epsilon, 82kD)
 integrin, beta 7
 integrin-linked kinase
 postmeiotic segregation increased 2-like 8
 ribosomal protein S6 kinase, 70kD, polypeptide 1
 gb:BC005688.1 /DEF=Homo sapiens. Similar to zinc finger protein 304, clone MGC:4079, mRNA, complete cds. /FEA=mRNA /PROD=Similar to zinc finger protein 304 /DB_XF
 KIAA0077 protein
 actin, gamma 1
 KIAA0618 gene product
 chromosome 19 open reading frame 5
 ubiquitin specific protease 1
 Consensus includes gb:BE878277 /FEA=EST /DB_XREF=gi:10327053 /DB_XREF=est:601488188F1 /CLONE=IMAGE:3890584 /UG=Hs.57079 Homo sapiens cDNA: FLJ1326
 basic leucine-zipper protein BZAP45
 PP3111 protein
 ribonuclease H1, large subunit
 ribosomal protein S17
 RecD protein-like (DNA helicase Q1-like)
 zinc finger protein 273
 Consensus includes gb:AL365505 /DEF=Human DNA sequence from clone RP11-382A12 on chromosome 20 Contains ESTs, STSs, GSSs and CpG islands. Contains the 5 pr
 G protein pathway suppressor 1
 proteasome (prosome, macropain) subunit, alpha type, 1
 DKF2PS47E1010 protein
 CD74 antigen (invariant polypeptide of major histocompatibility complex, class II antigen-associated)
 KIAA0100 gene product
 Consensus includes gb:AK025759.1 /DEF=Homo sapiens cDNA: FLJ22106 fls, clone HEP17675. /FEA=mRNA /DB_XREF=gi:10438374 /UG=Hs.323470 Homo sapiens cDNA: AK025759
 major histocompatibility complex, class I, A
 cyclin A2
 annexin A2
 mitochondrial import receptor Tom22
 suppressor of Ty 4 homolog 1 (S. cerevisiae)
 GrpE-like protein co-chaperone

NM_001985
 BC004118
 A0215854
 A424797
 NM_002817
 NM_003074
 M92301
 AF085736
 NM_006936
 NM_014390
 AL031228
 AV702627
 NM_014754
 AL050205
 AL050205
 A4747426
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 AJ001626
 AF062463
 BF000409
 NM_001905
 NM_019056
 AF130103
 NM_014730
 NM_006002
 NM_025164
 AF152863
 BC000478
 NM_000560
 BC003360
 A1582285
 NM_024546
 NM_013411
 U23028
 NM_000889
 NM_004517
 A1375694
 M60725
 BC005868
 D38521
 BC001920
 AK022555
 NM_018174
 NM_003368
 BE878277
 NM_014670
 NM_022156
 NM_006397
 BC004886
 NM_002907
 AU132789
 AL365505
 NM_004127
 BC005932
 AF261137
 K01144
 NM_014680
 AK025759
 A923492
 NM_001237
 NM_004039
 NM_020243
 BC002802
 AL542571

glutamate receptor, ionotropic, AMPA 3	NM_000828	1.87
coatomer protein complex, subunit alpha	U24105	1.87
GABA(A) receptor-associated protein	NM_007278	1.86
synaptic nuclei-expressed gene 1b	AF043290	1.86
H1 histone family, member 4	AL353759	1.86
chromosome 11 open reading frame 9	AC004770	1.86
S-phase kinase-associated protein 2 (p45)	BC001441	1.86
transcription factor Op-1	R50866	1.86
similar to Bos tauus P14 protein	NM_030877	1.86
hypothetical protein FLJ13962	NM_024862	1.86
jumping translocation breakpoint	AF151058	1.86
leptin receptor overlapping transcript-like 1	AF161461	1.86
actin, beta	X00351	1.86
BTG3 associated nuclear protein	NM_017869	1.86
acid phosphatase 1, soluble	BE872974	1.86
nucleophosmin/nucleoplamin 3	NM_006993	1.86
osteoclast stimulating factor 1	NM_012383	1.86
RAB6A, member RAS oncogene family	NM_002869	1.86
epilepsy, progressive myoclonus type 2, Lafora disease (laforin)	BC005286	1.86
SVI/SMF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	D26156	1.86
actin, beta	AA809056	1.86
basic leucine zipper transcription factor, ATF-like	AC007182	1.86
aquaporin 7	NM_001170	1.86
ribosomal protein L10	AW057781	1.86
Consensus includes gb:18Q249599 /FEA=EST /DB_XREF=gi:12759426 /DB_XREF=est:602319692F1 /CLONE=IMAGE:4414895 /UG=Hs.110771 Homo sapiens cDNA: FLJ211	BQ249599	1.85
ataxia telangiectasia and Rad3 related	U49844	1.85
citrate synthase	BC000105	1.85
actin, gamma 1	NM_001614	1.85
CGI-86 protein	AL022316	1.85
dynactin 3 (p22)	NM_007234	1.85
poly(A) binding protein, cytoplasmic 1	AI734929	1.85
hypothetical protein FLJ10036	NM_017875	1.85
PAP-1 binding protein	NM_031288	1.85
talin 1	NM_006289	1.85
ring finger protein 14	NM_004290	1.85
hypothetical protein FLJ12542	NM_024899	1.85
KIAA0542 gene product	AA521233	1.85
homolog of yeast RNase MRP/RNase P protein Pop5	NM_015918	1.85
cleavage and polyadenylation specific factor 1, 160kD subunit	NM_013291	1.84
poly(A) binding protein, nuclear 1	NM_004643	1.84
nucleoside phosphorylase	NM_000270	1.84
cortistatin	AB000263	1.84
glucose regulated protein, S8kD	DB3485	1.84
procollagen (type III) N-endopeptidase	NM_002768	1.84
RNA binding motif protein 14	NM_006328	1.84
casein kinase 1, delta	NM_001893	1.84
H4 histone family, member D	NM_003541	1.84
suppressor of Ty, 5 homolog (S. cerevisiae)	NM_003169	1.84
MMS19-like (MET18 homolog, S. cerevisiae)	NM_022362	1.84
CDC14 cell division cycle 14 homolog A (S. cerevisiae)	AF064103	1.84
Consensus includes gb:AL354872 /DEF=Human DNA sequence from clone RP11-42O15 on chromosome 1. Contains ESTs, STSs, GSSs and a CpG island. Contains the CTH	AL354872	1.84
KIAA0699 protein	BC002327	1.84
eukaryotic translation initiation factor 2, subunit 1 (alpha, 35kD)	BC002513	1.84
protein kinase, cAMP-dependent, catalytic, beta	NM_002731	1.84
chaperonin containing TCP1, subunit 6A (zeta 1)	NM_001762	1.84
ligatin	NM_006893	1.84
RAP2B, member of RAS oncogene family	NM_002886	1.84
quinoid dihydropteridine reductase	BC000376	1.84
mitochondrial ribosomal protein L24	NM_024540	1.84
transmembrane protein 4	NM_014255	1.83
reticulon 2	AF004222	1.83
homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1	AF217990	1.83
phorbol-12-myristate-13-acetate-induced protein 1	NM_021127	1.83

actin, beta	NM_001101	1.83
tyrosine kinase, non-receptor, 1	NM_003985	1.83
hypothetical protein FLJ13812	BC002449	1.83
acidic protein rich in leucines	AV712577	1.83
ribonuclease H1	BG534527	1.83
immunoglobulin heavy constant mu	BG340670	1.83
SET translocation (myeloid leukemia-associated)	D45188	1.83
protease (prosome, macropain) subunit, alpha type, 1	NM_002786	1.83
hypothetical protein FLJ20897	AI613383	1.83
carbonyl reductase	NM_016286	1.83
DKFZP566C0424 protein	BG171020	1.82
non-metastatic cells 2, protein (NM23B) expressed in	NM_002512	1.82
Consensus includes gb N92498 FEA-EST DB_XREF=gi:264807 DB_XREF=est:zb28a04.s1 CLONE=IMAGE:304878 UG=Hs.326248 Homo sapiens cDNA: FLJ22071 fs.	N92498	1.82
cleavage and polyadenylation specific factor 5, 25 kD subunit	NM_007006	1.82
hypothetical protein FLJ10769	NM_018210	1.82
jumping translocation breakpoint	NM_006694	1.82
signal transducer and activator of transcription 3 (acute-phase response factor)	AA634272	1.82
metallothionein 1H	NM_005951	1.82
RNA isopentenylpyrophosphate transferase	BE964125	1.82
HSPC156 protein	NM_014178	1.82
high-mobility group (nonhistone chromosomal) protein 17	BC003689	1.82
cell membrane glycoprotein, 110000M(r) (surface antigen)	NM_007002	1.82
KIAA0052 gene product	AA918224	1.82
CREBBP/EP300 inhibitory protein 1	AF109873	1.82
KIAA0071 protein	D31888	1.82
ATP citrate lyase	A1971281	1.82
Cip1-interacting zinc finger protein	AA054734	1.82
v-erb avian sarcoma virus CT10 oncogene homolog	AW612311	1.82
hypothetical protein MGC3731	NM_024313	1.82
DNA segment on chromosome 21 (unique) 2056 expressed sequence	NM_003683	1.82
RuvB-like 1 (E. coli)	BC000519	1.82
adaptor-related protein complex 2, beta 1 subunit	AL567295	1.82
eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD)	AL031313	1.82
adenomatosis polyposis coli	A1375486	1.81
arginine-glutamic acid dipeptide (RE) repeats	AB036737	1.81
putative membrane protein	AF277194	1.81
basic transcription factor 3	X74070	1.81
ribosomal protein S10	AA320784	1.81
Consensus includes gb AL133228 DEF=Human DNA sequence from clone RPS-1071L10 on chromosome 20 Contains part of a gene for a new member of the thymosinlike	AL133228	1.81
protease (prosome, macropain) 26S subunit, non-ATPase, 7 (Mov34 homolog)	NM_002811	1.81
ADP-ribosylation factor 5	NM_001662	1.81
protein kinase C, iota	L18964	1.81
vacuolar protein sorting factor 4A	NM_013245	1.81
UDP-glucose pyrophosphorylase 2	NM_006759	1.81
araA (bacterial) arsenite transporter, ATP-binding, homolog 1	NM_004317	1.81
hypothetical protein FLJ11183	NM_018351	1.81
KIAA0811 protein	NM_014944	1.81
hypothetical protein FLJ20530	NM_017864	1.81
KIAA1012 protein	NM_014839	1.80
KIAA0217 protein	AU154358	1.80
staufen, RNA binding protein (Drosophila)	BC000830	1.80
heterogeneous nuclear ribonucleoprotein A2/B1	NM_002137	1.80
protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcineurin A gamma)	NM_005605	1.80
cyclic-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4)	U36945	1.80
high-mobility group (nonhistone chromosomal) protein 17-like 3	NM_006353	1.80
KIAA0759 protein	AB018302	1.80
glutamy-prolyl-RNA synthetase	AI475965	1.80
angle-associated, migratory cell protein	NM_001087	1.80
glutathione-S-transferase like; glutathione transferase omega	NM_004832	1.80
hypothetical protein DKFZ056A00523	AL527334	1.80
hypothetical protein FLJ11583	AA388062	1.80
midustin 1	U95622	1.80
CGI-45 protein	NM_015999	1.80

eukaryotic translation termination factor 1	M75715	1.80
CpG binding protein	NM_014593	1.79
erbB2 interacting protein	NM_018895	1.79
hypothetical protein FLJ11198	NM_018358	1.79
putative receptor protein	NM_003876	1.79
DNA segment on chromosome X (unique) 9879 expressed sequence	NM_006014	1.79
RecD protein-like (DNA helicase Q1-like)	BC001052	1.79
DKF ZP564D116 protein	BE962354	1.79
chromosome 8 open reading frame 11	BC000388	1.79
CGI-38 protein	NM_015963	1.79
nuclear receptor subfamily 1, group D, member 1	X72631	1.79
COP9 homolog	BC003090	1.79
inhibitor of DNA binding 3, dominant negative helix-loop-helix protein	NM_002167	1.79
SET translocation (myeloid leukemia-associated)	NM_015959	1.79
leucine-rich PPR-motif containing	A276616	1.79
HLA class II region expressed gene KE2	M92439	1.79
myosin IE	NM_014280	1.79
chromosome 11 hypothetical protein ORF3	NM_004998	1.79
butyrophilin, subfamily 3, member A2	NM_020154	1.79
ribosomal protein L17	BC002832	1.79
splicing factor 3b, subunit 2, 145KD	NM_000985	1.79
KIAA0217 protein	NM_008842	1.79
sorting nexin 1	AW628886	1.79
phosphoglycerate dehydrogenase	NM_003099	1.79
MIL1 protein	NM_008623	1.78
polydromus cofactor synthesis 2	W84482	1.78
isocitrate dehydrogenase 3 (NADP+) beta	NM_015367	1.78
putative protein similar to nussy (Drosophila)	NM_004531	1.78
protein kinase C, nu	NM_006899	1.78
short-chain dehydrogenase/reductase 1	NM_005768	1.78
ligase I, DNA, ATP-dependent	NM_005813	1.78
TF53 target gene 1	NM_004753	1.78
non-canonical ubiquitin conjugating enzyme 1	NM_000234	1.78
uncharacterized bone marrow protein BM042	AB007458	1.78
KIAA0522 protein	AF151039	1.78
nuclear receptor subfamily 5, group A, member 2	NM_018456	1.78
peroxisomal membrane protein 2 (22KD)	BF571239	1.78
hypothetical protein FLJ10287	AF146343	1.78
aryl hydrocarbon receptor interacting protein	NM_018663	1.78
bromodomain-containing 4	NM_018049	1.78
MCM4 minichromosome maintenance deficient 4 (S. cerevisiae)	NM_003977	1.78
hypothetical protein MGC10963	BF718610	1.78
brefeldin A-inhibited guanine nucleotide-exchange protein 1	AA904621	1.78
hypothetical protein DKF Zp564K0822	NM_030573	1.77
uncharacterized hematopoietic stem/progenitor cells protein MDS032	AK025637	1.77
single-stranded DNA-binding protein 1	NM_030796	1.77
UDP-Gal-beta-GlcNAc beta 1,4-galactosyltransferase, polypeptide 4	NM_018467	1.77
gp.NM_014128.1 /DEF-Homo sapiens PRO0470 protein (PRO0470), mRNA, /FEA-mRNA (GEN=PRO0470) PROD=PRO0470 protein /DB_XREF=gj:7662555 /UG=Hs.278937	NM_003143	1.77
KIAA0573 protein	BC004523	1.77
leucine-rich PPR-motif containing	NM_014128	1.77
tubulin tyrosine ligase-like 1	BC006374	1.77
heterogeneous nuclear ribonucleoprotein C (C1/C2)	A653608	1.77
nuclear transcription factor Y, gamma	NM_012263	1.77
guanine nucleotide binding protein (G protein), beta polypeptide 1	AV725195	1.77
hypothetical protein	BC005003	1.77
complement component 1, q subcomponent binding protein	AF070603	1.77
hypothetical protein MGC5244,	BF038386	1.77
DnaJ (Hsp40) homolog, subfamily B, member 12	AU151801	1.77
hypothetical protein	NM_031213	1.77
SCAN domain-containing 1	NM_017626	1.77
zinc finger protein 259, pseudogene	NM_015705	1.77
	NM_016558	1.77
	Z95118	1.77

Interferon, alpha 21	M12350	1.77
aryl hydrocarbon receptor interacting protein	AL556532	1.77
Integrin, alpha 4 (antigen CD49D, alpha 4 subunit of VLA-4 receptor)	NM_000885	1.77
arginine-rich, mutated in early stage tumors	NM_006010	1.77
tubulin-interacting protein	AL080147	1.77
ATPase, aminophospholipid transporter (APLT), Class I, type 8A, member 1	AB013452	1.77
hypothetical protein FLJ22104	AK000684	1.77
lipase A, lysosomal acid, cholesterol esterase (Wolman disease)	NM_000235	1.77
pumilio homolog 1 (Drosophila)	BG474429	1.76
X-ray repair complementing defective repair in Chinese hamster cells 5 (double-strand-break rejoining; Ku autoantigen, 80kD)	AA205834	1.76
HCR (a-helix coiled-coil rod homologue)	AF589868	1.76
APGS autophagy 5-like (S. cerevisiae)	AF293841	1.76
hypothetical protein FLJ21908	NM_024604	1.76
tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation protein, zeta polypeptide	NM_003406	1.76
phosphoprotein regulated by mitogenic pathways	NM_025195	1.76
leucine-rich and death domain containing	AF274972	1.76
SH3-domain, GRB2-like, endophilin B2	AW131863	1.76
glutathione synthetase	L42531	1.76
cytoplasmic polyadenylation element binding protein	NM_030594	1.76
mitochondrial ribosomal protein L42	BE782148	1.76
regulator of G-protein signalling 10	AI744627	1.76
cyclin G2	L49506	1.76
hypothetical protein DKFZp434G0522	NM_017566	1.76
res-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1)	BG292367	1.76
phosphomevalonate kinase	NM_008556	1.76
small acidic protein	NM_014267	1.76
LAT1-3TM protein	NM_031211	1.76
BCS1-like (yeast)	NM_004328	1.76
DKFZP564J0123 protein	BC002873	1.75
sorbitol dehydrogenase	L29008	1.75
pituitary tumor-transforming 1 interacting protein	NM_004339	1.75
thymosin, beta 10	NM_021103	1.75
serine protease inhibitor, Kunitz type, 2	AF027205	1.75
hypothetical protein FLJ14040	AK024102	1.75
signal recognition particle 72kD	AF069765	1.75
homolog of Yeast RRP4 (ribosomal RNA processing 4), 3'-5'-exoribonuclease	NM_014285	1.75
transcriptional adaptor 3 (ADA3, yeast homolog)-like (PCAF histone acetylase complex)	AK024982	1.75
protein phosphatase 4 (formerly X), catalytic subunit	BC001416	1.75
ribosomal protein S3A	NM_001006	1.75
DKFZP586O0120 protein	NM_014077	1.75
H3 histone family, member H	NM_003534	1.75
clone FLB4739	AA133341	1.75
Consensus includes gb:AF064606.1 /DEF=Homo sapiens KB07 protein mRNA, partial cds. /FEA=mRNA /PROD=KB07 protein /DB_XREF=gi:3152665 /UG=Hs.306242 Homo	NM_005065	1.75
sef-1 suppressor of lin-12-like (C. elegans)	AL060112	1.75
Consensus includes gb:AL080112.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H0722 (from clone DKFZp586H0722). /FEA=mRNA /DB_XREF=gi:5262539 /UG=Hs.33273	NM_000112	1.75
matrin 3	NM_018834	1.75
SET translocation (myeloid leukemia-associated)	AV702810	1.75
hypothetical protein FLJ13194	NM_025146	1.75
gb:Z25421.1 /DEF=H sapiens protein-serine/threonine kinase gene, complete cds. /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405720 /FL=gb.Z25421	NM_025146	1.75
CDC-like kinase 2	Z25421	1.75
casein kinase 1, delta	NM_003993	1.75
ribosomal protein S17	AV700224	1.75
collin	NM_001021	1.75
postmeiotic segregation increased 2-like 11	NM_004645	1.75
DNA segment on chromosome X (unique) 9923 expressed sequence	U38980	1.75
hypothetical protein FLJ20244	NM_004699	1.75
KIAA0194 protein	BC002492	1.75
DKFZP547E2110 protein	D83778	1.75
cytochrome c oxidase subunit VIIa polypeptide 2 (liver)	AB028628	1.75
KIAA0731 protein	AC004544	1.75
outer dense fibre of sperm tails 2	BE881529	1.75
GLE1 RNA export mediator-like (yeast)	AF053970	1.75
HSPC009 protein	NM_001489	1.75
	NM_014019	1.75

numb homolog (Drosophila)	AF015040	1.75
vascular endothelial growth factor	AF022375	1.75
ribosomal protein L36a	NM_001001	1.75
protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcineurin B, type I)	NM_000845	1.75
lactate dehydrogenase C	NM_002301	1.75
serine/threonine kinase 19	NM_004197	1.75
ES1 (zebrafish) protein, human homolog of	NM_004649	1.75
guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3	J03198	1.74
hypothetical protein F23985	BF115644	1.74
hypothetical protein	NM_016404	1.74
nif endonuclease III-like 1 (E. coli)	U79718	1.74
NAD(P) dependent steroid dehydrogenase-like; H105e3	BC000245	1.74
membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10)	NM_007287	1.74
H2B histone family, member A	BC001131	1.74
RING1 and YY1 binding protein	AB029551	1.74
nuclear transcription factor, X-box binding 1	AW291398	1.74
N-sulfoglucosamine sulfohydrolase (sulfamidase)	U30894	1.74
adaptor-related protein complex 2, alpha 2 subunit	AI25280	1.74
RPB5-mediated protein	AB006572	1.74
DKFZP434D1335 protein	AL117499	1.74
ribosomal protein L12	NM_000976	1.74
pseudouridylyl synthase 1	NM_025215	1.74
phosphatidylinositol 4-kinase, catalytic, alpha polypeptide	NM_002650	1.74
G-2 and S-phase expressed 1	BF106380	1.74
culin 4B	AF650819	1.74
methylene tetrahydrofolate dehydrogenase (NAD+ dependent), methenyltetrahydrofolate cyclohydrolase	NM_006638	1.73
G protein-coupled receptor 68	U35398	1.73
paired-like homeodomain transcription factor 1	NM_002653	1.73
Consensus includes gb:AU144066 /FEA=EST /DB_XREF=est:AU144066 /CLONE=HEMBA1000798 /UG=Hs.285519 Homo sapiens cDNA FLJ111904	AU144066	1.73
sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1	NM_001055	1.73
HSPC015 protein	NM_014026	1.73
tumor necrosis factor receptor superfamily, member 6b, decoy	U04810	1.73
muS homolog 3 (E. coli)	NM_004226	1.73
serine/threonine kinase 17b (apoptosis-inducing)	AW245400	1.73
tyrosyl-RNA synthetase	U63139	1.73
RAD50 homolog (S. cerevisiae)	AI882255	1.73
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 9kD	NM_021960	1.73
myeloid cell leukemia sequence 1 (BCL2-related)	NM_005778	1.73
RNA binding motif protein 5	BE544663	1.73
Consensus includes gb:BE544663 /FEA=EST /DB_XREF=gi:9773308 /DB_XREF=est:601074131F1 /CLONE=IMAGE:3460398 /UG=Hs.110407 ESTs, Weakly similar to coded	BC000120	1.73
general transcription factor IIF, polypeptide 1 (74kD subunit)	BF971587	1.73
tubulin, beta polypeptide	AI195589	1.73
hypothetical protein DKFZP434E1723	NM_012234	1.73
RING1 and YY1 binding protein	NM_002389	1.73
membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen)	AU135134	1.73
a disintegrin and metalloproteinase domain 10	O25304	1.73
Rac/Cdc42 guanine exchange factor (GEF) 6	NM_020123	1.73
SM-11044 binding protein	AA827892	1.72
Cluster incl. AA827892:cd08e04.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:1387358 /clone_end=3' /gb=AA827892 /gi=2900255 /ug=Hs.239016 /len=622	NM_016062	1.72
CD112 protein	BC001604	1.72
hematopoietic protein 1	NM_001030	1.72
ribosomal protein S27 (metalloproteinin 1)	AB018272	1.72
KIAA0729 protein	NM_018131	1.72
hypothetical protein FLJ10540	NM_002583	1.72
PRKC, apoptosis, WT1, regulator	NM_004271	1.72
MD-1, RP105-associated	NM_006808	1.72
protein translocation complex beta	L23928	1.72
serine hydroxymethyltransferase 1 (soluble)	AA628948	1.72
adenylosuccinate synthase	U37012	1.72
cleavage and polyadenylation specific factor 1, 160kD subunit	NM_003720	1.72
Down syndrome critical region gene 2	S60876	1.72
zinc finger protein, subfamily 1A, 1 (IkaroS)	AW025150	1.72
mitogen-activated protein kinase kinase kinase 12		

putative nuclear protein
 CG1-146 protein
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4
 putative neurotransmitter receptor
 proteasome (prosome, macropain) subunit, beta type, 2
 ribosomal protein L30
 gb-Z25430.1 /DEF=H sapiens protein-serine/threonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405738 /FL=gb-Z25430. Z25430
 MAP/microtubule affinity-regulating kinase 3
 methionine-tRNA synthetase
 methylrhodanese phosphorylase
 oxoglutarate (alpha-ketoglutarate) dehydrogenase (lipoamide)
 small nuclear ribonucleoprotein D1 polypeptide (16kD)
 lorcin
 EAP30 subunit of ELL complex
 R0D1 regulator of differentiation 1 (S. pombe)
 L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain
 hypothetical protein hCLA-iso
 likely ortholog of mouse tubulin alpha 4
 transportin-SR
 Consensus includes gb:AK001816.1 /DEF=Homo sapiens cDNA FLJ10854 fts, clone PLACE1000383, highly similar to Homo sapiens mRNA for MTMR1 protein. /FEA=mRNA
 alcohol dehydrogenase 5 (class III), chi polypeptide
 ATP citrate lyase
 tumor suppressor deleted in oral cancer-related 1
 chromosome 14 open reading frame 2
 protein tyrosine phosphatase, receptor type, N
 hypothetical protein FLJ22301
 Interleukin 4 receptor
 proteasome (prosome, macropain) activator subunit 1 (PA28 alpha)
 cadherin and integrin binding protein (DNA-dependent protein kinase interacting protein)
 gb:NM_006328.1 /DEF=Homo sapiens seven transmembrane domain protein (NIFE14), mRNA. /FEA=mRNA /GEN=NIFE14 /PROD=seven transmembrane domain protein /C
 B-cell CLL/lymphoma 6 (zinc finger protein 51)
 myosin, light polypeptide 4, alkali; atrial, embryonic
 KIAA0785 protein
 NADH dehydrogenase (ubiquinone) Fe-S protein 5 (15kD) [NADH-coenzyme Q reductase]
 cytochrome c oxidase subunit VIa polypeptide 1
 gb:AF116705.1 /DEF=Homo sapiens PRO2489 mRNA, complete cds. /FEA=mRNA /PROD=PRO2489 /DB_XREF=gi:7959908 /FL=gb:AF116705.1
 M-phase phosphoprotein 9
 polymerase (DNA-directed) sigma
 ataxia telangiectasia mutated (Includes complementation groups A, C and D)
 small nuclear RNA activating complex, polypeptide 3, 50kD
 arachne homolog 2 (Drosophila)
 transformin-2 alpha (htr-2 alpha)
 Cluster Incl. AA150503.208g04.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-481788 /clone_end=3 /gb=AA150503 /gi=1722017 /ug=Hs.24435 /len=631
 dihydrolipoamide dehydrogenase (E3 component of pyruvate dehydrogenase complex, 2-oxo-glutarate complex, branched chain keto acid dehydrogenase complex)
 bromodomain-containing 7
 hypothetical protein FLJ14393
 actin related protein 2/3 complex, subunit 1B (41 kD)
 non-POU-domain-containing, octamer-binding
 aquaporin 3
 STE20-like kinase
 eukaryotic translation initiation factor 2B, subunit 3 (gamma, 58kD)
 gb:NM_015870.1 /DEF=Homo sapiens endogenous retrovirus H D1 leader region/integrase-derived ORF1, ORF2, and putative envelope protein (HSU88895), mRNA. /FEA=ml
 SWAP-70 protein
 Cluster Incl. AA628140.297e06.s1 Homo sapiens cDNA, 3' end /clone=IMAGE-1055746 /clone_end=3 /gb=AA628140 /gi=2540527 /ug=Hs.126089 /len=302
 Kalsio
 small fragment nuclease
 mitochondrial ribosomal protein S18B
 hypothetical protein MGC2491
 protein disulfide isomerase-related protein
 similar to S. cerevisiae SSMA4
 phosphatidylserine receptor
 ubiquinol-cytochrome c reductase (6.4kD) subunit
 chloride channel, nucleotide-sensitive, 1A

nuclease sensitive element binding protein 1	BC002411	1.69
general transcription factor IIF, polypeptide 1 (74kD subunit)	AW190445	1.69
hypothetical protein	NM_019005	1.69
P450 (cytochrome) oxidoreductase	AF258341	1.69
S100 calcium binding protein A4 (calcium protein, calvasculin, metastasin, murine placental homolog)	NM_002961	1.69
ELKS protein	AB029004	1.69
hypothetical protein	NM_015932	1.69
hypothetical protein FLJ13984	NM_024770	1.69
ras homolog gene family, member G (rho G)	NM_001665	1.69
ribosomal protein L7	NM_000971	1.69
sialidase 1 (lysosomal sialidase)	U84246	1.69
KIAA0746 protein	AB018289	1.69
coatamer protein complex, subunit alpha	AB021079	1.69
hypothetical protein FLJ22940	NM_024571	1.69
transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C)	NM_005648	1.69
pM5 protein	N39536	1.69
sortilin-related receptor, L (OLR class) A repeats-containing	NM_003105	1.69
hypothetical protein MGC2668	NM_030914	1.68
ribosomal protein L21	NM_000982	1.68
glycosyltransferase AD-017	NM_018446	1.68
RAN binding protein 3	AB032261	1.68
stearyl-CoA desaturase (delta-9-desaturase)	AB032261	1.68
HSPCQ38 protein	NM_025084	1.68
ADP-ribosylation factor related protein 1	NM_016096	1.68
promyelocytic leukemia	AF230401	1.68
hypothetical protein MGC10753	NM_003224	1.68
galactokinase 2	NM_016628	1.68
hypothetical protein FLJ11773	NM_002044	1.68
U2(RNU2) small nuclear RNA auxiliary factor 1 (non-standard symbol)	NM_021934	1.68
embryonic ectoderm development	NM_008758	1.68
hypothetical protein FLJ11896	AF080227	1.68
hypothetical protein MGC2731	NM_024611	1.68
mitochondrial carrier homolog 1	NM_024068	1.68
proteasome (prosome, macropain) subunit, beta type, 6	AF189289	1.68
cytochrome c oxidase subunit VIb pseudogene-3	BC000835	1.68
KIAA0999 protein	AL031594	1.67
Consensus includes gb:N95443 /FEA=EST /DB_XREF=gi:1267753 /DB_XREF=est:zb81c12.s1 /CLONE=IMAGE-310006 /UG=Hs.19.180 Homo sapiens mRNA; cDNA DKFZp5: N95443	AA044154	1.67
B-cell receptor-associated protein BAP29	NM_018844	1.67
S-adenosylmethionine decarboxylase 1	M21154	1.67
KIAA0871 protein	AF112221	1.67
Consensus includes gb-AV708130 /FEA=EST /DB_XREF=gi:107253395 /DB_XREF=est:AV708130 /CLONE=ADCAMA05 /UG=Hs.283965 Homo sapiens scavenger receptor cti	AF143684	1.67
myosin IXB	BC005212	1.67
hypothetical protein FLJ20003	NM_013366	1.67
anaphase-promoting complex subunit 2	BC005161	1.67
hypothetical protein MGC4638	NM_005340	1.67
histidine triad nucleotide binding protein	NM_023005	1.67
bromodomain adjacent to zinc finger domain, 1B	M18003	1.67
ferredoxin 1	BC002585	1.67
RAB7, member RAS oncogene family-like 1	NM_025233	1.67
nucleotide binding protein	S72931	1.67
c8.1A	NM_014418	1.67
T-cell leukemia/lymphoma 6	NM_006469	1.67
NS1-binding protein	NM_014183	1.67
HSPC162 protein	AL553254	1.66
DEADH (Asp-Glu-Ala-Asp-His) box polypeptide 19 (DBP5 homolog, yeast)	U47077	1.66
protein kinase, DNA-activated, catalytic polypeptide	NM_005507	1.66
cofilin 1 (non-muscle)	U56331	1.66
sarcoglycan, delta (35kD dystrophin-associated glycoprotein)	AF080157	1.66
conserved helix-loop-helix ubiquitous kinase	BC006479	1.66
hypothetical protein MGC2749	AB051480	1.66
hypothetical protein FLJ20719	AI186701	1.66
inhibitor of growth family, member 1-like		

nuclease sensitive element binding protein 1	X96666	1.66
ubiquitin fusion degradation 1-like	BC001049	1.66
ribosomal protein L13	BC004954	1.66
proline synthetase co-transcribed (bacterial homolog)	AL138616	1.66
G-rich RNA sequence binding factor 1	BF058465	1.66
ORF	NM_015853	1.66
uncoupling hydrogen decarboxylase	AW614435	1.66
19A24 protein	NM_021181	1.66
ribosomal protein L7	BG389744	1.66
hypothetical protein	AA463853	1.66
hyaluronan-mediated motility receptor (RHAMM)	U29343	1.66
chromatin assembly factor 1, subunit A (p150)	BF062223	1.66
hypothetical protein FLJ10975	NM_018288	1.66
KIAA0052 protein	D29841	1.66
spermine synthase	NM_004595	1.66
amplified in osteosarcoma	AI653900	1.66
ubiquitin specific protease 22	AA621731	1.66
hypothetical protein FLJ12587	AL136796	1.66
EH-domain containing 4	AF320070	1.66
platelet-activating factor acetylhydrolase, isoform 1b, alpha subunit (45kD)	L13387	1.66
hypothetical protein FLJ11021 similar to splicing factor, arginine/serine-rich 4	BE398879	1.66
ATP-binding cassette, sub-family C (CFTR/MRP), member 9	NM_005691	1.66
Consensus includes gb:AK028168.1 /DEF=Homo sapiens cDNA; FLJ22515 fls, clone HRC12122, highly similar to AF052101 Homo sapiens clone 23872 mRNA sequence.	AF028168	1.66
SNARE protein	NM_006555	1.65
KIAA1118 protein	AB029041	1.65
keratohelin (impodin) beta 3	AU148466	1.65
hypothetical protein FLJ22116	NM_024624	1.65
gene from NF2 meningioma region of 22q12	BC003815	1.65
suppression of tumorigenicity 7	NM_018412	1.65
ELK1, member of ETS oncogene family	M25269	1.65
RAP1B, member of RAS oncogene family	NM_015646	1.65
polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)	X62006	1.65
Rho guanine nucleotide exchange factor (GEF) 7	NM_003899	1.65
Consensus includes gb:AU144378 /FEA=EST /DB_XREF=gi:11005899 DB_XREF=est:AU144378 /CLONE=HEMBA1001734 /UG=Hs.296650 Homo sapiens cDNA FLJ11475	AU144378	1.65
Cytidine-rich protein 1 (intestinal)	NM_001311	1.65
SEC14-like 1 (S. cerevisiae)	NM_003003	1.65
zinc finger protein 217	NM_006526	1.65
RAD17 homolog (S. pombe)	AF096533	1.65
transcriptional intermediary factor 1	AL538264	1.65
HSPC159 protein	NM_014181	1.65
mitochondrial ribosomal protein L35	NM_016622	1.65
CGI-125 protein	NM_016080	1.65
hypothetical protein FLJ20756	NM_017952	1.65
KIAA0247 gene product	NM_014734	1.65
protein kinase D2	AL050147	1.65
DAZ associated protein 1	NM_018959	1.65
RAB3 GTPase-ACTIVATING PROTEIN	AI040009	1.65
ATPase, H ⁺ transporting lysosomal (vacuolar proton pump), member M	AF100741	1.65
leucine-rich repeat-containing 2	NM_024512	1.65
chondroitin sulfate proteoglycan 6 (barnacan)	AI373676	1.65
zinc finger protein 216	AW471220	1.65
nuclear distributor gene C (A. nidulans) homolog	AF241788	1.65
aldo-keto reductase family 7, member A3 (aflatoxin aldehyde reductase)	NM_012067	1.65
protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (lprn), alpha 1	U22815	1.65
ceroid-lipofuscinosis, neuronal 5	AV700514	1.65
protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (lprn), alpha 1	BG033593	1.65
hypothetical protein R0083M05.2	NM_021167	1.64
elastase 3A, pancreatic (protease E)	BC005918	1.64
growth factor, augmentor of liver regeneration (ERV1 homolog, S. cerevisiae)	AF124604	1.64
protease, serine, 25	NM_013247	1.64
branched chain aminotransferase 2, mitochondrial	NM_001190	1.64
low density lipoprotein receptor defect C complementing	NM_007357	1.64
MAX protein	AA723514	1.64

purinergic receptor P2X, ligand-gated ion channel, 2
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 5 (RNA helicase, 58kD)
 microfilament-associated protein 2
 hypothetical protein MGC11256
 Consensus includes gb:AW304232 /FEA=EST /DB_XREF=gi:6713921 /DB_XREF=ast:vx82g01.x1 /CLONE=IMAGE:2825040 /UG=Hs.181357 laminin receptor 1 (67kD, riboso
 DNA (cytosine-5)-methyltransferase 1
 hypothetical protein FLJ10083
 histidine triad nucleotide binding protein
 inhibitor of growth family, member 3
 elastase 3A, pancreatic (protease E)
 transcription factor A, mitochondrial
 eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD)
 ring finger protein 13
 hypothetical protein MGC4614
 somatostatin receptor 5
 interleukin enhancer binding factor 3, 90kD
 eukaryotic translation elongation factor 1 epsilon 1
 CGI-96 protein
 histidine triad nucleotide binding protein
 JTV1 gene
 polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)
 hypothetical protein DC42
 exosome component Rrp41
 H2A histone family, member M
 transducer of ERBB2, 1
 actin, gamma 1
 hypothetical protein FLJ10439
 Consensus includes gb:AL049250.1 /DEF=Homo sapiens mRNA; cDNA DKFZ564D113 (from clone DKFZ564D113) /FEA=mRNA /DB_XREF=gi:4499989 /UG=Hs.194637
 regulator of G-protein signalling 14
 chromosome 9 open reading frame 12
 asf2 (absent, small, or homeotic)-like (Drosophila)
 hypothetical protein
 metastasis-associated 1-like 1
 hypothetical protein FLJ11021 similar to splicing factor, arginine/serine-rich 4
 ribosomal protein L12
 hypothetical protein FLJ12547
 hypothetical gene MGC19595
 phospholipase C, gamma 1 (formerly subtype 14b)
 prohibitin
 Consensus includes gb:AK025862.1 /DEF=Homo sapiens cDNA: FLJ22209 fs, clone HRC01495 /FEA=mRNA /DB_XREF=gi:10438505 /UG=Hs.288528 Homo sapiens cDNA
 zinc finger protein 79 (pT7)
 tumor necrosis factor, alpha-induced protein 2
 integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)
 zinc finger protein 42 (myeloid-specific retinoic acid-responsive)
 huntingtin interacting protein-1-related
 hypothetical protein FLJ12671
 eukaryotic translation initiation factor 4A, isoform 2
 aldolase C, fructose-bisphosphate
 JTV1 gene
 differentially expressed in FDCP 6 homolog (mouse)
 DKFZ564J157 protein
 putative dimethyladenosine transferase
 protein kinase C substrate 80K-H
 MAGUK protein p55T, Protein Associated with Lins 2
 hypothetical protein
 hypothetical protein PP2447
 mitogen-activated protein kinase phosphatase x
 dihydropyrimidine S-acetyltransferase (E2 component of pyruvate dehydrogenase complex)
 proline arginine-rich end leucine-rich repeat protein
 KIAA1028 protein
 FK506 binding protein 4 (59kD)
 arafane homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila)

NM_016318 1.64
 NM_004398 1.64
 NM_017459 1.64
 NM_024324 1.64
 AW304232 1.64
 NM_001379 1.64
 NM_017992 1.64
 U27143 1.64
 NM_019071 1.64
 O00308 1.64
 NM_012251 1.64
 BC002719 1.64
 AF070558 1.64
 NM_024294 1.64
 NM_001053 1.64
 BG032366 1.64
 NM_004280 1.64
 AL022316 1.64
 M33197 1.64
 N32864 1.64
 AI828526 1.64
 BC004383 1.64
 NM_030921 1.64
 AL039469 1.63
 NM_003513 1.63
 AA675892 1.63
 BE741683 1.63
 NM_018093 1.63
 NM_049250 1.63
 E037195 1.63
 NM_022755 1.63
 AB020982 1.63
 BF038366 1.63
 BE222618 1.63
 NM_023012 1.63
 AK026491 1.63
 NM_024992 1.63
 W67887 1.63
 AL022394 1.63
 AL560017 1.63
 AK025862 1.63
 X65232 1.63
 AI882445 1.63
 A4215654 1.63
 NM_003422 1.63
 AB014555 1.63
 NM_030980 1.63
 NM_001987 1.63
 NM_005165 1.63
 NM_006303 1.63
 NM_022047 1.63
 NM_018457 1.63
 BC002841 1.63
 NM_002743 1.63
 NM_016447 1.63
 AL096741 1.63
 BG399562 1.63
 NM_020185 1.63
 BF978872 1.62
 U41344 1.62
 AB028951 1.62
 AA894574 1.62
 NM_005744 1.62

gb-NM_013307.1 /DEF=Homo sapiens non-functional folate binding protein (HSAF000381). mRNA. /FEA=CDS /GEN=HSAF000381 /PROD=non-functional folate binding prot NM_013307
 Consensus includes gb:AL049761 /DEF=Human DNA sequence from clone RP5-863C7 on chromosome 20p12.3-13. Contains the CSNK2A1 gene for casein kinase 2 alpha 1 AL049761
 enoyl Coenzyme A hydratase, short chain, 1, mitochondrial
 adaptor-related protein complex 2, alpha 2 subunit
 KIAA0737 gene product
 hypothetical protein
 zinc finger protein 288
 lactate dehydrogenase A
 MAD, mothers against decapentaplegic homolog 4 (Drosophila)
 UDP glycosyltransferase 1 family, polypeptide A1
 mitotic spindle coiled-coil related protein
 CD22 antigen
 polymorphisiderodermis autoantigen 1 (75kD)
 calcium binding protein Cab45 precursor
 DKFZP364G2022 protein
 EH-domain containing 1
 integrin cytoplasmic domain-associated protein 1
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 3
 Consensus includes gb:A062978 /FEA=EST /DB_XREF=gi:5755693 /DB_XREF=est:W25605.x1 /CLONE=IMAGE:2508513 /UG=Hs 288908 WAS protein family, member 2 /FL A062978
 brefeldin A-inhibited guanine nucleotide-exchange protein 2
 ubiquitin-conjugating enzyme E2G 2 (UBC7 homolog, yeast)
 tubulin, alpha 1 (testis specific)
 similar to MRJ gene for a member of the DNAP protein family (H. sapiens)
 high-mobility group (nonhistone chromosomal) protein 17-like 3
 HSPC128 protein
 CDC45 cell division cycle 45-like (S. cerevisiae)
 hypothetical protein FLJ20038
 hypothetical protein FLJ23027
 zinc finger protein 161 homolog (mouse)
 DKFZP34J154 protein
 DKFZP368D0623 protein
 pinin, desmosome associated protein
 phosphoribosyl pyrophosphatase synthetase-associated protein 2
 v-rel reticuloendotheliosis viral oncogene homolog B, nuclear factor of kappa light polypeptide gene enhancer in B-cells 3 (avian)
 nuclear factor (erythroid-derived 2)-like 1
 tubulin-specific chaperone e
 ribosomal protein S24
 carnitine palmitoyltransferase I, muscle
 TED protein
 origin recognition complex, subunit 3-like (yeast)
 eukaryotic translation initiation factor 1A
 DKFZP586G011 protein
 E51 (zebrafish) protein, human homolog of
 sperm associated antigen 4
 Consensus includes gb:BE733579 /FEA=EST /DB_XREF=gi:10147971 /DB_XREF=est:601571007F1 /CLONE=IMAGE:3925800 /UG=Hs 82202 ribosomal protein L17
 intercellular adhesion molecule 2
 suppression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein)
 macrophage myristoylated alanine-rich C kinase substrate
 ubiquitination factor E4B (UFD3 homolog, yeast)
 similar to S. cerevisiae Sec6p and R. norvegicus rsec6
 ubiquinol-cytochrome c reductase complex (7.2 kD)
 transcription factor A, mitochondrial
 Consensus includes gb:AC007204 /DEF=Homo sapiens chromosome 19, BAC 2732339 (CIT-B-320G13) /FEA=CDS /DB_XREF=gi:4559317 /UG=Hs 284229 Homo sapiens chr AC007204
 CCR4-NOT transcription complex, subunit 3
 Consensus includes gb:AJ224869 /DEF=Homo sapiens CXCR4 /FEA=mRNA /DB_XREF=gi:3059119 /UG=Hs 89414 chemokine (C-X-C motif) AJ224869
 cylin-dependent kinase 4
 KIAA0542 gene product
 staufen, RNA binding protein (Drosophila)
 growth arrest and DNA-damage-inducible, beta
 N-acetyltransferase, homolog of S. cerevisiae ARO1
 Consensus includes gb:AL021395 /DEF=Human DNA sequence from clone RP1-269M15 on chromosome 20q12.1-13.12 Contains a gene similar to peptidylprolyl isomerase (cy AL021395)
 MAD, mothers against decapentaplegic homolog 2 (Drosophila)
 WW domain binding protein 2
 U79458

KIAA1069 protein
formin-binding protein 3
chaperonin containing TCP1, subunit 8B (zeta 2)
C3H-type zinc finger protein; similar to D. melanogaster muscibind B protein
pro-oncogene receptor inducing membrane injury gene
serine/threonine kinase 38
squamous cell carcinoma antigen recognised by T cells
chromosome 21 open reading frame 4
polymerase (DNA directed), delta 1, catalytic subunit (125kD)
Consensus includes gb:BE568184 /FEA=EST /DB_XREF=gi:9811904 /DB_XREF=est1601341722F1 /CLONE=IMAGE:3683860 /UG=Hs.200400 Human DNA sequence from B
general transcription factor IIE, polypeptide 2 (beta subunit, 34kD)
ribosomal protein S4, X-linked
Friend leukemia virus integration 1
HSPC028 protein
synaptoglycin 2
cat eye syndrome chromosome region, candidate 5
Consensus includes gb:A1972416 /FEA=EST /DB_XREF=gi:5769332 /DB_XREF=est:wr39610.x1 /CLONE=IMAGE:2490067 /UG=Hs.24385 Human hbc647 mRNA sequence
APG12 autophagy 12-like (S. cerevisiae)
chromosome 11 hypothetical protein ORF4
hydroxacyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit
mitogen-activated protein kinase kinase 7
hypothetical protein FLJ10631
Consensus includes gb:A1687738 /FEA=EST /DB_XREF=gi:4899032 /DB_XREF=est:tp93g08.x1 /CLONE=IMAGE:2206910 /UG=Hs.699 peptidylprolyl isomerase B (cyclophilin A)
ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast)
activating transcription factor 4 (tax-responsive enhancer element B67)
H2A histone family, member 2
plasminogen activator, urokinase receptor
metallothionein 1H
putative membrane protein
chromosome 14 open reading frame 1
Consensus includes gb:BC003542.1 /DEF=Homo sapiens, clone IMAGE:3611719, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3611719) /DB_XREF=BC003542
POPA (processing of precursor, S. cerevisiae) homolog
gb:M87268.1 /DEF=human Igm VDJ-region mRNA, complete cds. /FEA=CDS /GEN=IGHM /PROD=igm /DB_XREF=gi:186197 /FL=gb:M87268.1
KIAA0399 protein
tubulin, alpha, ubiquitous
DKFZP434B168 protein
Sec61 gamma
LIN-7 protein 3
KIAA0293 protein
ornithine decarboxylase antizyme inhibitor
nucleix (nucleoside diphosphate linked moiety X)-type motif 6
hypothetical protein FLJ12768
hypothetical protein
myosin regulatory light chain interacting protein
v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian)
splicing factor, arginine/serine-rich 2
benzodiazepine receptor (peripheral)
ribosomal protein L13
hypothetical protein FLJ12492
protein similar to E.coli yhdg and R. capsulatus nlr3
natriuretic peptide precursor A
mitogen-activated protein kinase kinase 5
hypothetical protein FLJ10359
fumarate hydratase
dihydropyrimidine S-succinyltransferase (E2 component of 2-oxo-glutarate complex)
hypothetical protein FLJ10377
sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4C
Cluster Ind. A1363375:q53f12.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2015951 /done_end=3 /gb=A1363375 /gi=4114996 /ug=Hs.179801 /len=470
c-src tyrosine kinase
thyroid hormone receptor interactor 8
sirtuin silent mating type information regulation 2 homolog 7 (S. cerevisiae)
c-Mpl binding protein
ribosomal protein S6

v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neurofiblioblastoma derived oncogene homolog (avian)
 C3HC4-type zinc finger protein
 translational inhibitor protein p14.5
 ubiquitin specific protease 12
 EZIG2 protein
 KIAA0240 protein
 eukaryotic translation initiation factor 2, subunit 1 (alpha, 35kD)
 similar to Caenorhabditis elegans protein C42C1.9
 glutamine-fructose-6-phosphate transaminase 1
 v-Ha-ras Harvey rat sarcoma viral oncogene homolog
 KIAA0170 gene product
 coronin, actin binding protein, 1C
 eukaryotic translation initiation factor 4B
 hypothetical protein
 Golgi apparatus protein 1
 protein kinase, AMP-activated, gamma 1 non-catalytic subunit
 Treacher Collins-Franceschetti syndrome 1
 hypothetical protein FLJ20808
 CGI-51 protein
 zinc finger protein 262
 hypothetical protein MGC3062
 BRCA1-interacting protein 1
 chondroitin sulfate proteoglycan BEHAB/brevican
 guanidinoacetate N-methyltransferase
 coxsackie virus and adenovirus receptor
 polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein I)
 SMC2 structural maintenance of chromosomes 2-like 1 (yeast)
 adaptor-related protein complex 2, alpha 2 subunit
 phosphotyrosinotid transfer protein
 inositol(myo)-1(or 4)-monophosphatase 2
 hypothetical protein MGC5509
 solute carrier family 25 (mitochondrial carrier, citrate transporter), member 1
 hypothetical protein PRO2730
 phosphoinositide 3-kinase, platelet
 solute carrier family 21 (organic anion transporter), member 8
 nucleosome assembly protein 1-like 1
 deam-associated protein
 neighbor of COX4
 KIAA0409 protein
 KIAA0182 protein
 Consensus includes gb:AL181956.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761A17121 (from clone DKFZp761A17121). /FEA=mRNA /DB_XREF=gi:7328008 /UG=Hs.306-
 BCL2-like 1
 binder of A1 Two
 interferon, alpha 13
 KIAA0462 protein
 hepatoma-derived growth factor (high-mobility group protein 1-like)
 transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)
 eukaryotic translation initiation factor 2B, subunit 1 (alpha, 26kD)
 tetrapeptide repeat domain 1
 ribosomal protein L12
 zinc finger protein 36, C3H type-like 2
 Cbort10 protein
 KIAA0555 gene product
 eukaryotic translation initiation factor 3, subunit 9 (eta, 116kD)
 serpin-specific protease
 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 10 (42kD)
 kinesin-associated protein 3
 HSPC125 protein
 hematopoietic PBX-interacting protein
 exonuclease 1
 Consensus includes gb:BG332462 /FEA=EST /DB_XREF=gi:13138900 /DB_XREF=est:602432837F1 /CLONE=IMAGE-4550505 /UG=Hs.3343 phosphoglycerate dehydrogenase
 hypothetical protein FLJ10099
 fatty-acid-Coenzyme A ligase, long-chain 1

X03363
 NM_024835
 N54448
 AF022790
 NM_016565
 AB01951
 NM_004084
 AL568449
 BE645771
 NM_005343
 AB73553
 BC002342
 NM_001417
 NM_019048
 AK025457
 NM_002733
 NM_000356
 NM_017960
 NM_015380
 AB50586
 NM_024065
 AF360549
 NM_021948
 NM_001156
 NM_001338
 BF690062
 NM_008444
 BC006155
 NM_006224
 NM_014214
 NM_024053
 U25147
 AL137654
 NM_002627
 NM_019844
 AB85751
 NM_004394
 NM_006067
 NM_015324
 AA206181
 AL161956
 NM_001191
 BF244411
 NM_006900
 AB007831
 NM_004494
 AA768906
 NM_001414
 NM_003314
 AA281332
 A356398
 BE963765
 NM_014750
 U78525
 NM_021627
 NM_004544
 NM_014970
 NM_014165
 AB35162
 NM_003686
 BG332462
 NM_017894
 NM_001995

hypothetical protein FLJ13491	AI431902	1.57
MAD1 mitotic arrest deficient-like 1 (yeast)	NM_003550	1.57
cytochrome c oxidase subunit VIc pseudogene 1	AF042165	1.57
cell division cycle 2-like 2	AL031282	1.57
hypothetical protein FLJ22490	BE044503	1.57
macrophage stimulating 1 (hepatocyte growth factor-like)	NM_020998	1.57
hydroxycyl glutathione hydrolase	NM_005326	1.57
KIAA0084 protein	D42043	1.57
peroxisomal short-chain alcohol dehydrogenase	NM_021004	1.57
CGI-18 protein	NM_015947	1.57
zinc finger protein 289, ID1 regulated	BE299871	1.57
thymopoleilin	BG391171	1.57
RAB7, member RAS oncogene family	BG281416	1.57
hypothetical protein MGC10763	AI560217	1.57
KIAA0943 protein	NM_013325	1.57
eukaryotic translation initiation factor 2, subunit 2 (beta, 38KD)	AL031668	1.57
UDP-Gal:betaGlcNAc beta 1,4-galactosyltransferase, polypeptide 6	AF097159	1.57
replication factor C (activator 1) 1 (145KD)	L14922	1.57
methyl-CpG binding domain protein 4	AL358619	1.57
Consensus includes gb:AV733266 /FEA=EST /DB_XREF=gi:10850811 /DB_XREF=est:AV733266 /CLONE=cdAAUG04 /UG=Hs.76325 step II splicing factor SLU7	AV733266	1.57
Inositol polyphosphate-5-phosphatase, 145kD	NM_005541	1.57
CGG triplet repeat binding protein 1	NM_003663	1.57
RAD50 homolog (S. cerevisiae)	NM_005732	1.57
hypothetical protein MGC5466	AI559507	1.57
gb:AF293339.1 /DEF=Homo sapiens ornithine decarboxylase antizyme 4 mRNA, complete cds. /FEA=mRNA /PROD=ornithine decarboxylase antizyme 4 /DB_XREF=gi:11024; /FEA=EST /DB_XREF=gi:12674986 /DB_XREF=est:602340822F /CLONE=IMAGE:4448789 /UG=Hs.82202 ribosomal protein L17	AF293339	1.57
fibroblast growth factor 4 (heparin secretory transforming protein 1, Kaposi sarcoma oncogene)	BF510715	1.57
nucleolin	NM_005381	1.57
isoprenylcysteine carboxyl methyltransferase	AL578502	1.57
low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor)	BF304759	1.57
KIAA0650 protein	AA868754	1.57
hypoxia-inducible factor 1, alpha subunit (basic helix-loop-helix transcription factor)	NM_001530	1.57
actin, beta	DS0604	1.57
splicing factor (CC1:3)	NM_004902	1.57
Consensus includes gb:G168283 /FEA=EST /DB_XREF=gi:12674986 /DB_XREF=est:602340822F /CLONE=IMAGE:4448789 /UG=Hs.82202 ribosomal protein L17	BG168283	1.57
N-acetylglucosamine kinase	NM_017567	1.57
KIAA0251 protein	AK025504	1.57
Ewing sarcoma breakpoint region 1	AF327066	1.57
X-box binding protein 1	NM_005080	1.57
huntingtin interacting protein 2	NM_005339	1.57
U2 small nuclear ribonucleoprotein auxiliary factor (65KD)	NM_007279	1.57
Consensus includes gb:AL050217.1 /DEF=Homo sapiens mRNA; cDNA DKFZp5860523 (from clone DKFZp5860523). /FEA=mRNA /DB_XREF=gi:4884458 /UG=Hs.308487	AL050217	1.57
steroid sulfatase (microsomal), arylsulfatase C, isozyme S	AU138166	1.57
hypothetical protein KIAA1164	AL049226	1.57
zinc finger protein 22 (K0X 15)	AA744771	1.57
RAN binding protein 7	AA939270	1.56
hypothetical protein FLJ12910	NM_024573	1.56
ATPase, (Na+ /K+)-transporting, beta 4 polypeptide	AW975818	1.56
DKFZp586C0424 protein	NM_012069	1.56
cytochrome c oxidase subunit VIIb pseudogene 1	AL050028	1.56
elF4 associated protein HSPC021	AF042164	1.56
isocitrate dehydrogenase 3 (NAD+) gamma	NM_016091	1.56
glutaryl-Coenzyme A dehydrogenase	U69268	1.56
Rab geranylgeranyltransferase, beta subunit	NM_000159	1.56
hypothetical protein BK1048E9.5	U49245	1.56
KIAA0874 protein	NM_022081	1.56
protection of telomeres 1	AW572809	1.56
thioredoxin interacting protein	BC002923	1.56
hypothetical protein DKFZp564O0523	AA812232	1.56
lactate dehydrogenase B	AL136619	1.56
golgi associated, gamma adaptin ear containing, ARF binding protein 1	NM_002300	1.56
vesicle-associated membrane protein 3 (cellubrevin)	AW009695	1.56
serine/threonine kinase 12	NM_004781	1.56
	AB011446	1.56

putative GTP-binding protein similar to RAY/RAB1C	NM_006860	1.56
proteasome (prosome, macropain) 26S subunit, non-ATPase, 13	NM_002817	1.56
nuclear transcription factor Y, alpha	AL031778	1.56
amyloid beta (A4) precursor-like protein 2	BC004371	1.56
cyclin B2	NM_004701	1.56
cytochrome c oxidase subunit IV isoform 1	AA854966	1.56
ubiquitin specific protease 10	BC000263	1.56
uncharacterized hematopoietic stem/progenitor cells protein MDS026	AI138993	1.56
protein kinase, interferon-inducible double stranded RNA dependent activator	AF083033	1.56
BRF1 homolog, subunit of RNA polymerase III transcription initiation factor IIB (S.cerevisiae)	N91109	1.56
ADP-ribosylation factor 3	M33384	1.56
nuclear localization signal deleted in velocardiofacial syndrome	NM_003776	1.56
putative L-type neutral amino acid transporter	AW000954	1.56
polymyositis/scleroderma autoantigen 2 (100kD)	NM_002885	1.56
matrix metalloproteinase 11 (stromelysin 3)	NM_005940	1.56
DKFZP564K2062 protein	NM_015421	1.56
allantoinase	NM_018436	1.55
hypothetical protein FLJ12615 similar to membrane protein, palmitoylated 3 (MAGUK p55 subfamily member 5)	NM_022474	1.55
protease, serine, 15	U02389	1.55
potassium voltage-gated channel, Shal-related subfamily, member 1	NM_004879	1.55
nascent-polypeptide-associated complex alpha polypeptide	NM_005594	1.55
keratin 6B	L42612	1.55
cytochrome c oxidase subunit Vb	AU557312	1.55
Mad4 homolog	AA631438	1.55
CDC5 cell division cycle 5-like (S. pombe)	AW268817	1.55
regulator of G-protein signalling 2, 24kD	NM_002923	1.55
hypothetical protein FLJ10808	NM_018227	1.55
protoporphyrinogen oxidase	NM_000309	1.55
transgelin 2	BC002616	1.55
regulator of G-protein signalling 10	NM_002925	1.55
KIAA0285 protein	AW205215	1.55
dashrin, light polypeptide (Lcb)	NM_001834	1.55
laminin, alpha 5	BC003355	1.55
translocase of inner mitochondrial membrane 17 homolog A (yeast)	BC004439	1.55
hypothetical protein MGC11279	NM_024326	1.55
LDL receptor adaptor protein	AA169780	1.55
tubulin, gamma 1	NM_001070	1.55
spermidine/spermine N1-acetyltransferase	BES71383	1.55
ClpX caseinolytic protease X homolog (E. coli)	NM_006660	1.55
hypothetical protein FLJ20568	AA438930	1.55
zinc finger protein 258	NM_007167	1.55
spectrin, alpha, non-erythrocytic 1 (alpha-fodrin)	U83867	1.55
hypothetical protein MGC20727	AU879381	1.55
macrophage stimulating 1 (hepatocyte growth factor-like)	U37055	1.55
cathepsin C	NM_001814	1.55
zinc finger protein 354A	NM_005849	1.55
palmitoyl-protein thioesterase 1 (ceroid-lipofuscinosis, neuronal 1, infantile)	NM_000310	1.55
stearoyl-CoA desaturase (delta-9-desaturase)	AF116816	1.55
alpha integrin binding protein 63	NM_007216	1.55
hepatic leukemia factor	M95565	1.55
BAF53	NM_004301	1.55
cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle)	NM_000079	1.55
A kinase (PRKA) anchor protein 13	M90360	1.55
tRNA-guanine transglycosylase	NM_031209	1.55
leucyl/cystinyl aminopeptidase	NM_005575	1.55
EH-domain containing 1	NM_006795	1.55
hypothetical protein	AF722190	1.54
phosphatidylinositol binding clathrin assembly protein	NM_024053	1.54
hypothetical protein MGC861	AU856493	1.54
dCMP deaminase	BG469030	1.54
putative mitochondrial outer membrane protein import receptor	D13891	1.54
inhibitor of DNA binding 2, dominant negative helix-loop-helix protein	X63381	1.54
MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)		

polycystic kidney disease 1 (autosomal dominant)	AK023376	1.54
capillary morphogenesis protein 1	AI610355	1.54
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1	NM_003079	1.54
uroporphyrinogen decarboxylase	M14016	1.54
pleckstrin homology-like domain, family A, member 1	AA576961	1.54
activating transcription factor 3	NM_001674	1.54
thioredoxin-like 2	AF118652	1.54
CD1A antigen, a polypeptide	M28625	1.54
PTD015 protein	NM_014040	1.54
NOL1R2 protein	AK021688	1.54
X-ray repair complementing defective repair in Chinese hamster cells 1	NM_006297	1.54
protein tyrosine phosphatase, receptor type, C	NM_002836	1.54
acyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase)	NM_001607	1.54
CGI-152 protein	NM_020410	1.54
nuclear factor (erythroid-derived 2)-like 2	NM_006164	1.54
GIM2 ganglioside activator protein	X62076	1.54
SMARTHDAC1 associated repressor protein	NM_015001	1.54
casein kinase 2, alpha 1 polypeptide	NM_001895	1.54
hypothetical protein FLJ12428	NM_022783	1.54
dynactin 4	AI038068	1.54
TCF3 (E2A) fusion partner (in childhood Leukemia)	NM_013342	1.54
Consensus includes gb:A742553 /FEA=EST /DB_XREF=gi:5110841 /DB_XREF=est:wg55d02.x1 /CLONE=IMAGE:2368995 /UG=Hs.184592 protein kinase, lysine deficient 1 /	AI742553	1.54
hypothetical protein FLJ10539	NM_018130	1.54
neuroblastoma RAS viral (v-ras) oncogene homolog	NM_002524	1.54
RNA binding motif protein 6A	AI738479	1.54
actin, gamma 1	BC026805	1.54
NADH dehydrogenase (ubiquinone) Fc-S protein 7 (20kD) (NADH-coenzyme Q reductase)	BC005954	1.54
formin-binding protein 17	AB011126	1.54
ubiquitously-expressed transcript	NM_004182	1.54
mitochondrial ribosomal protein L34	AB048652	1.53
copper chaperone for superoxide dismutase	NM_005125	1.53
superoxide dismutase 2, mitochondrial	W46388	1.53
homolog of yeast ribosome biogenesis regulatory protein RRS1	BC001811	1.53
ION3 protein	NM_015384	1.53
sialyltransferase 7D (alpha-N-acetylneuraminyl-2,3-beta-galactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase)	NM_014403	1.53
major histocompatibility complex, class I-like sequence	NM_001531	1.53
basigin (OK blood group)	AL550657	1.53
steroid dehydrogenase homolog	NM_016142	1.53
nuclear transcription factor Y, gamma	NM_014223	1.53
nucleolar protein GU2	AF334103	1.53
gb:BC002746.1 /DEF=Homo sapiens, Similar to dodecenoyl-Coenzyme A delta isomerase (3,2 trans-enoyl-Coenzyme A isomerase), clone MGC:3903, mRNA, complete cds. /	BC002746	1.53
hypothetical protein	AI081779	1.53
MAX protein	M64240	1.53
cyclin I	AF135162	1.53
ribosomal protein L14	AA638274	1.53
hypothetical protein FLJ20640	BC000890	1.53
programmed cell death 5	NM_004708	1.53
hypothetical protein FLJ10747	NM_018202	1.53
hypothetical protein FLJ20452	NM_017828	1.53
heterogeneous nuclear ribonucleoprotein K	BC000355	1.53
Consensus includes gb:AK022459.1 /DEF=Homo sapiens cDNA FLJ12397 /ts, clone MAMMA1002769, weakly similar to Homo sapiens cell cycle progression restoration 8 prot	AK022459	1.53
PDGFA associated protein 1	NM_014891	1.53
protein kinase, lysine deficient 1	AI445745	1.53
EDF transcription factor 1	M96577	1.53
RNA binding protein	AF251062	1.53
metallothionein 1F (functional)	M10943	1.53
hypothetical protein DKFZ0781H1710	NM_031297	1.53
Consensus includes gb:BE379006 /FEA=EST /DB_XREF=gi:5324371 /DB_XREF=est:801236272F1 /CLONE=IMAGE:3608680 /UG=Hs.99766 Homo sapiens mRNA; cDNA D	BE379006	1.53
malate dehydrogenase 2, NAD (mitochondrial)	BC001917	1.53
myogenic factor 3	BC000353	1.53
phosphatidylinositol 4-kinase, catalytic, beta polypeptide	NM_002651	1.53
CAMP responsive element binding protein 3 (human)	AF028674	1.53
MCM3 mitochondrial maintenance deficient 3 (S. cerevisiae) associated protein	AJ010089	1.53

potassium channel, subfamily K, member 13	NM_022054	1.53
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	NM_003070	1.53
diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)	M60276	1.53
serine/threonine kinase 19	NM_003274	1.52
transmembrane protein 1	NM_000928	1.52
phospholipase A2, group IB (pancreas)	M57732	1.52
transcription factor 1, hepatic; LF-B1, hepatic nuclear factor (HNF1), albumin proximal factor	NM_000994	1.52
ribosomal protein L32	AF191653	1.52
nudix (nucleoside diphosphate linked moiety X)-type motif 4	NM_004896	1.52
vacuolar protein sorting 28 (yeast)	BC000966	1.52
gamma tubulin ring complex protein (76p gene)	NM_004922	1.52
SEC24 related gene family, member C (S. cerevisiae)	D84105	1.52
membrane cofactor protein (CD46, trophoblast-lymphocyte cross-reactive antigen)	NM_021971	1.52
GDP-mannose pyrophosphorylase B	NM_003377	1.52
vascular endothelial growth factor B	BC003375	1.52
mitochondrial ribosomal protein L3	U62136	1.52
ubiquitin-conjugating enzyme E2 variant 2	NM_001023	1.52
ribosomal protein S20	NM_001023	1.52
Consensus includes gb:AL109722.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIIMAGE 31619 /FEA=mRNA /DB_XREF=gi:5689814 /UG=Hs.170079 Homo sapiens	U67280	1.52
calumenin	AF087853	1.52
growth arrest and DNA-damage-inducible, beta	NM_014847	1.52
KIAA0144 gene product	NM_001157	1.52
annexin A11	BC004467	1.52
KIAA0171 gene product	NM_016821	1.52
8-oxoguanine DNA glycosylase	NM_002629	1.52
phosphoglycerate mutase 1 (brain)	NM_004890	1.52
sperm associated antigen 7	M36172	1.52
myosin, light polypeptide 4, alkali, atrial, embryonic	NM_016172	1.52
putative glioblastoma cell differentiation-related	NM_004941	1.52
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 8 (RNA helicase)	NM_0022061	1.52
Consensus includes gb:AK022061.1 /DEF=Homo sapiens cDNA FLJ11989 fls. clone HEMBB1001527 /FEA=mRNA /DB_XREF=gi:10433381 /UG=Hs.306622 Homo sapiens	T79953	1.52
KIAA0040 gene product	D89846	1.52
nucleoside homologue 6 (E. coli)	NM_006247	1.52
protein phosphatase 3, catalytic subunit	AF336851	1.52
hypothetical protein FLJ20596	NM_025230	1.52
hypothetical protein PRO2389	NM_025230	1.52
nucleoporin 98KD	A4527238	1.52
zinc finger protein 22 (KOX 15)	NM_006963	1.52
myotubularin related protein 3	AF233437	1.52
peptidylprolyl isomerase F (cyclophilin F)	BC003080	1.52
HSPC1171 protein	NM_000833	1.52
glutamate receptor, ionotropic, N-methyl D-aspartate 2A	NM_014296	1.52
calpain 7	NM_000661	1.52
ribosomal protein L9	AC004531	1.52
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 28	NM_006363	1.51
Sec23 homolog B (S. cerevisiae)	NM_002248	1.51
potassium intermediate/small conductance calcium-activated channel, subfamily N, member 1	NM_021993	1.51
FUS interacting protein (serine-arginine rich) 2	NM_005332	1.51
hemoglobin, zeta	NM_001707	1.51
B-cell CLL/lymphoma 7B	NM_030940	1.51
hypothetical protein MGC4276 similar to CG8198	AL562282	1.51
hypothetical protein PP591	D87453	1.51
mitochondrial ribosomal protein S27	NM_022471	1.51
hypothetical protein FLJ13057 similar to germ cell-less	N36997	1.51
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 16	AK000145	1.51
Consensus includes gb:AK000145.1 /DEF=Homo sapiens cDNA FLJ20138 fls. clone COL07172 /FEA=mRNA /DB_XREF=gi:7020041 /UG=Hs.306384 Homo sapiens	NM_013239	1.51
protein phosphatase 2A 48 kDa regulatory subunit	M90355	1.51
basic transcription factor 3, like 2	AF006080	1.51
growth hormone 2	AU154985	1.51
adenylyate kinase 2	BF026595	1.51
ribosomal protein S17	AF172452	1.51
opioid growth factor receptor	AJ236912	1.51
ADP-ribosyltransferase (NAD+-poly(ADP-ribose) polymerase)-like 2		

CGI-100 protein	NM_016040	1.51
Ins-related tyrosine kinase 3 ligand	NM_001459	1.51
hypothetical protein	NM_018988	1.51
esterase Diformylglutathione hydrolase	BC001169	1.51
hypothetical protein	NM_013298	1.51
phosphorylase kinase, beta	BG149218	1.51
cyclin B1	BE407516	1.51
NCK adaptor protein 1	NM_006153	1.51
spectrin, alpha, non-erythrocytic 1 (alpha-fodrin)	AL110273	1.51
hypothetical protein, clone Telhont(italy_B41)_Strat02270_FL142	NM_020247	1.51
truncaloid repeat containing 12	AL563727	1.51
X-ray repair complementing defective repair in Chinese hamster cells 4	AB017445	1.51
v-rel reticuloendotheliosis viral oncogene homolog (avian)	NM_002908	1.51
phospholipase A2-activating protein	AF145020	1.51
hypothetical protein FLJ20558	NM_017880	1.51
plative membrane protein	AB020980	1.51
chaperonin containing TCP1, subunit 2 (beta)	NM_006431	1.51
heterogeneous nuclear ribonucleoprotein A3	BG505670	1.51
hypothetical protein PRO1843	BF247371	1.51
CGI-48 protein	NM_016001	1.51
hypothetical protein MGC4054	NM_024341	1.51
CGI-07 protein	NM_015938	1.51
KIAA0415 gene product	AB007675	1.51
Consensus includes gb:W95043 /FEA=EST /DB_XREF=gi:1424182 /DB_XREF=est:zh46c11.s1 /CLONE=IMAGE:415124 /UG=Hs.79732 fibulin 1	W95043	1.51
FK506 binding protein 4 (S9KD)	NM_002014	1.51
G protein beta subunit-like	NM_022372	1.51
hypothetical protein FLJ22408	NM_024794	1.51
calreticulin	NM_004343	1.51
KIAA0476 gene product	NM_014856	1.51
hypothetical protein KIAA1164	AU121431	1.51
protein tyrosine phosphatase, non-receptor type 11	L07527	1.51
mitochondrial translational release factor 1	NM_004294	1.51
tumor protein 63 kDa with strong homology to p53	NM_003722	1.51
Wolf-Hirschhorn syndrome candidate 2	AF101434	1.51
YY1 transcription factor	Z14077	1.51
CCCTC-binding factor (zinc finger protein)	NM_006565	1.51
glutaminyl-peptide cyclotransferase (glutaminyl cyclase)	NM_012413	1.50
Consensus includes gb:AL515874 /FEA=EST /DB_XREF=gi:12779367 /DB_XREF=est:AL515874 /CLONE=CSODA001.YF06 (3 prime) /UG=Hs.83086 Homo sapiens GT212 m	AL515874	1.50
protein phosphatase 2, regulatory subunit B (B56), delta isoform	AB000835	1.50
ubiquitin B	NM_018955	1.50
Consensus includes gb:A1348009 /FEA=EST /DB_XREF=gi:4085215 /DB_XREF=est:qp61b09.x1 /CLONE=IMAGE:1927481 /UG=Hs.12421 hypothetical protein	A1348009	1.50
cystathionase (cystathionine gamma-lyase)	NM_001902	1.50
Consensus includes gb:BE671941 /FEA=EST /DB_XREF=gi:10032405 /DB_XREF=est:7a45g02.x1 /CLONE=IMAGE:3221810 /UG=Hs.157078 Homo sapiens cDNA FLJ127932 BE671941	BE671941	1.50
CGI-105 protein	NM_016044	1.50
Interleukin 2 receptor, gamma (severe combined immunodeficiency)	NM_000206	1.50
hypothetical protein FLJ22427	AK026080	1.50
hypothetical protein FLJ21820	AK021672	1.50
peroxiredoxin 2	L19185	1.50
pyrophosphatase (inorganic)	NM_021129	1.50
laminin, alpha 3 (nicotin (150KD), kalinin (165KD), epiligrin)	X77598	1.50
golgi associated, gamma adaptin ear containing, ARF binding protein 2	BE646414	1.50
BCL2-associated athanogene 2	AF095192	1.50
16.7kd protein	NM_016139	1.50
hypothetical protein FLJ10154	NM_016011	1.50
hypothetical protein FLJ22570	NM_024872	1.50
Consensus includes gb:AL096828 /DEF=Human DNA sequence from clone RPS-963E22 on chromosome 20 Contains the 3 end of a novel gene similar to NY-REN-2 Antigen, general transcription factor IIF, polypeptide 1 (74kD subunit)	AL096828	1.50
vaiy-RNA synthetase 2	NM_002096	1.50
hypothetical protein FLJ21613 similar to rat corneal wound healing related protein	BE790854	1.50
adaptor-related protein complex 1, beta 1 subunit	NM_021929	1.50
nucleosome assembly protein 1-like 1	NM_001127	1.50
hypothetical protein FLJ23320	AW148801	1.50
X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound	NM_024872	1.50
	AL023653	1.50

ribosomal protein S27a	NM_002954	1.50
dynamitin 1-like	NM_012062	1.50
HGF activator	NM_001528	1.50
small acidic protein	BE748698	1.50
KIAA0193 gene product	NM_014766	1.50
mitochondrial ribosomal protein L46	NM_022163	1.50
hypothetical protein FLJ10547	AL048795	1.50
colony stimulating factor 1 (macrophage)	M37435	1.50
nuclear receptor coactivator 3	U80737	1.50
DKFZP586J0119 protein	AJ011307	1.50
breast cancer 2, early onset	NM_000059	1.49
integrin, alpha X (antigen CD11C (p150), alpha polypeptide)	M81695	1.49
melastasin associated 1	BC006177	1.49
gongli associated, gamma adaptin ear containing, ARF binding protein 1	NM_013365	1.49
chromosome 5 open reading frame 6	NM_016605	1.49
ATP-binding cassette, sub-family B (MDR/TPAP), member 8	NM_007188	1.49
E74-like factor 1 (ets domain transcription factor)	M82882	1.49
O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase)	NM_003605	1.49
hypothetical protein FLJ13114	H27225	1.49
M27830 Human 28S ribosomal RNA gene, complete cds (S, M, 3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M27830	1.49
KIAA0801 gene product	NM_014829	1.49
immediate early response 5	NM_016545	1.49
sorting nexin 17	NM_014748	1.49
capping protein (actin filament) muscle Z-line, alpha 2	AV655920	1.49
partner of RAC1 (arfpapin 2)	NM_012402	1.49
scratch homolog 1, zinc finger protein (Drosophila)	NM_031309	1.49
mitochondrial ribosomal protein S28	NM_014018	1.49
baculoviral IAP repeat-containing 5 (survivin)	NM_001188	1.49
H1 histone family, member 3	NM_005320	1.49
nucleolar protein (KKED repeat)	BE796327	1.49
L-3-hydroxycy-Coenzyme A dehydrogenase, short chain	NM_005327	1.49
aminolevulinatase, delta-, synthase 1	NM_006688	1.49
RAS p21 protein activator (GTPase activating protein) 1	M23812	1.49
hypothetical protein FLJ10307	NM_018053	1.49
2,4-dienoyl CoA reductase 1, mitochondrial	NM_001359	1.49
chromobox homolog 3 (HP1 gamma homolog, Drosophila)	BE748755	1.49
tumor protein p53-binding protein	AI150117	1.49
anaphase-promoting complex 1: meiotic checkpoint regulator	NM_022682	1.49
hypothetical protein	RS1077	1.49
hypothetical protein	NM_017571	1.49
H1 histone family, member 0	BC000145	1.49
hypothetical protein FLJ10450	NM_018095	1.49
retinoid acid receptor, alpha	T91506	1.49
sialyltransferase 7D ((alpha-N-acetylneuraminyl-2,3-beta-galactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase)	AB035172	1.49
tumor necrosis factor (ligand) superfamily, member 8	NM_001244	1.49
Consensus includes gb:AV681533 /FEA=EST /DB_XREF=gi:9882547 /DB_XREF=est:AV681533 /CLONE=GLCGTD12 /UG=Hs.293637 ESTs	AV681533	1.49
dysactin 4 (p82)	NM_016221	1.49
Consensus includes gb:BE547674 /FEA=EST /DB_XREF=gi:9776319 /DB_XREF=est:601076255F1 /CLONE=IMAGE:3462325 /UG=Hs.204169 ESTs	BE547674	1.49
TH1-like (Drosophila)	NM_016397	1.49
sperm associated antigen 11	AF168619	1.49
nuclear factor, interleukin 3 regulated	NM_005384	1.49
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson-W.	AB030077	1.49
hypothetical protein FLJ22637	NM_025165	1.49
progesterone receptor membrane component 1	AL547946	1.49
topoisomerase (DNA) III beta	D87012	1.49
serologically defined colon cancer antigen 8	U79271	1.48
serine/arginine repetitive matrix 2	X92110	1.48
HCGVII-1 protein	AB014609	1.48
endocytic receptor (macrophage mannose receptor family)	NM_002883	1.48
Ran GTPase activating protein 1	NM_002776	1.48
prosaposin (variant Gaucher disease and variant metachromatic leukodystrophy)	BC002506	1.48
programmed cell death 10	H38643	1.48
RAN binding protein 16		

putative mitochondrial outer membrane protein import receptor
golgin-67
three prime repair exonuclease 1
protein phosphatase 2 (formerly 2A), regulatory subunit B" (PR 72), alpha isoform and (PR 130), beta isoform
gb:AF081735.1 /DEF=Homo sapiens clone 023e08 MY032 protein mRNA, complete cds. /FEA=mRNA /PROD=My032 protein /DB_XREF=gi:12002005 /UG=Hs.307099 Homo
BRCA2 and CDKN1A interacting protein
NM_016567
hypothetical protein FLJ20477
PEST-containing nuclear protein
pleckstrin homology, Sec7 and coiled/coil domains 2 (cytohesin-2)
zinc finger RNA binding protein
delta-like 3 (Drosophila)
sialoadhesin
H3 histone family, member F
Consensus includes gb:AK023938.1 /DEF=Homo sapiens cDNA FLJ13676 fls, clone T1HYO1001401 /FEA=mRNA /DB_XREF=gi:10436033 /UG=Hs.287604 Homo sapiens c
KIAA0475 gene product
putative translation initiation factor
integrin, alpha L (trigden CD11A (p180), lymphocyte function-associated antigen 1; alpha polypeptide)
clone HQ0477 PRO0477p
RecQ protein-like 4
CpG binding protein
ornithine decarboxylase antizyme 2
deoxrin
chromosome 11 hypothetical protein ORF4
KIAA0971 protein
asparaginyl-RNA synthetase
methyl-CpG binding domain protein 2
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae)
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit g
putative nuclear RNA helicase
hypothetical protein FLJ22530
tubulin-interacting protein
hypothetical protein FLJ13902
H2A histone family, member O
tubulin, alpha, ubiquitous
IDN3 protein
KIAA0317 gene product
homolog of rat nadrin
Consensus includes gb:AL137403.1 /DEF=Homo sapiens mRNA: cDNA DKF Zp434L092 (from clone DKF Zp434L092) /FEA=mRNA /DB_XREF=gi:6807951 /UG=Hs.306457 H AL137403
calcium channel, voltage-dependent, L type, alpha 1D subunit
Consensus includes gb:BF115739 /FEA=EST /DB_XREF=gi:10985215 /DB_XREF=est:7n64008.x1 /CLONE=IMAGE:3589246 /UG=Hs.15740 Homo sapiens mRNA: cDNA DK BF115739
heterogeneous nuclear ribonucleoprotein A3
Consensus includes gb:AW139802 /FEA=EST /DB_XREF=gi:5143220 /DB_XREF=est:U1-H-B11-aeq-e-08-0-UI.s1 /CLONE=IMAGE:2770347 /UG=Hs.173108 Homo sapiens cDNA
hypothetical protein FLJ21016
Consensus includes gb:AL031602 /DEF=Human DNA sequence from clone RPS-1174N9 on chromosome 1p34.1-35.3. Contains the gene for a novel protein with IBR domain,
fizzled homolog 3 (Drosophila)
centromere protein A (17kD)
golgi SNAP receptor complex member 1
GTP cyclohydrolase 1 (topa-responsive dystonia)
proteasome (prosome, macropain) 26S subunit, non-ATPase, 1
secretory carrier membrane protein 2
zinc finger protein 216
secreted protein of unknown function
cytochrome c oxidase subunit VIIa polypeptide 2 like
C-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine:polypeptide-N-acetylglucosaminyl transferase)
DKFZP434J1813 protein
WD repeat domain 1
protein regulator of cytokinesis 1
signal recognition particle 72kD
interferon-related developmental regulator 2
Wilms' tumour 1-associated protein
hypothetical protein 394D8_6
BRCA1 associated RING domain 1
myosin, heavy polypeptide 9, non-muscle

VAMP (vesicle-associated membrane protein)-associated protein A (33kD)	AF154847	1.47
gene amplified in squamous cell carcinoma 1	AB037901	1.47
major histocompatibility complex, class I, F	A168379	1.47
pre-mRNA splicing factor 17	NM_015891	1.47
Consensus includes gb:AF131823.1 /DEF=Homo sapiens clone 24997 mRNA sequence, /FEA=mRNA /DB_XREF=gi:4408659 /UG=Hs.308951 Homo sapiens clone 24997 mF	AF131823	1.47
murY homolog (E. coli)	NM_012222	1.47
single Ig IL-1R-related molecule	NM_021805	1.47
thioredoxin-like 2	AL138831	1.47
Consensus includes gb:AL031669 /DEF=Human DNA sequence from clone RPA-68120 on chromosome 20q11.23-12 Contains a gene for a protein similar to RPL12 (ribosomal protein L12)	AL031669	1.47
cathepsin, light polypeptide (Lcb)	NM_007097	1.47
Consensus includes gb:AW452623 /FEA=EST /DB_XREF=gi:5993399 /DB_XREF=est UI-H-B13-ali-d-09-0-U1.s1 /CLONE=IMAGE:3088608 /UG=Hs.14846 Homo sapiens mR	AW452623	1.47
tubulin, alpha, ubiquitous	BC008481	1.47
glycogen synthase kinase 3 alpha	L40027	1.47
SH3-domain GRB2-like 1	NM_003025	1.47
hypothetical protein FLJ23018	NM_024810	1.47
LIM domain kinase 2	A118402	1.47
high-mobility group (nonhistone chromosomal) protein isoforms 1 and Y	AL117466	1.47
chloride channel, calcium activated, family member 3	NM_002131	1.47
COI-45 protein	NM_004921	1.47
casein kinase 1, alpha 1	NM_016013	1.47
SH2-B homolog	BF341845	1.47
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 7A	AF227988	1.47
sympkin; Huntingtin interacting protein 1	Z97056	1.47
developmentally regulated GTP binding protein 1	NM_004819	1.47
zinc finger protein with interaction domain	NM_004147	1.47
cullin 4B	NM_006626	1.47
lysine 3-monoxygenase/tyrosine 5-monoxygenase activation protein, beta polypeptide	NM_003588	1.47
Consensus includes gb:AI344141 /FEA=EST /DB_XREF=gi:4081347 /DB_XREF=est tc02d09.x1 /CLONE=IMAGE:2062673 /UG=Hs.229850 ESTs. Highly similar to G01251 R	AI344141	1.47
protein disulfide isomerase-related protein	NM_003742	1.47
adaptor-related protein complex 3, beta 1 subunit	NM_003664	1.47
zinc metalloproteinase (STE24 homolog, yeast)	NM_005857	1.47
hexokinase 1	NM_000188	1.47
B06 antigen	NM_016579	1.47
2-5'-oligoadenylate synthetase 2 (69-71 kD)	NM_016817	1.47
cell division cycle 2-like 2	NM_015976	1.47
hypothetical protein FLJ20445	D13640	1.47
Ras homolog enriched in brain 2	AL031282	1.47
thyroid hormone receptor interactor 12	NM_017824	1.47
Era G-protein-like 1 (E. coli)	NM_005614	1.47
purinergic receptor P2X, ligand-gated ion channel, 7	NM_004238	1.47
TNF receptor-associated factor 4	AL562733	1.47
KIAA0678 protein	NM_002562	1.47
Cluster ind. AL050297; Homo sapiens mRNA; cDNA DKFp564N123 (from clone DKFp564N123) /cds=UNKNOWN /gb=AL050297 /lg=4886448 /ug=Hs.25894 /len=1340	NM_004295	1.47
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 5	AB014578	1.47
KIAA0332 protein	AL050287	1.47
sorting nexin 3	NM_006837	1.47
vascular endothelial growth factor	A1184562	1.46
zinc finger protein 131 (clone pTZ-10)	AB047360	1.46
neighbor of A-kinase anchoring protein 95	H95344	1.46
G protein-coupled receptor 87	AY966301	1.46
nucleoporin 155kD	NM_014371	1.46
secretory carrier membrane protein 1	NM_023915	1.46
ADP37 protein	NM_004298	1.46
ADP-ribosylation factor-like 1	AF005037	1.46
putative T1/S12 receptor binding protein	AF260335	1.46
KIAA0246 protein	BE890745	1.46
U5 snRNP-specific 40 kDa protein (hPrp8-binding)	NM_006858	1.46
nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, beta	NM_015136	1.46
heat shock 10kD protein 1 (chaperonin 10)	AL157420	1.46
	A1684894	1.46
	NM_002157	1.46

hypothetical protein 394D8_6
excision repair cross-complementing rodent repair deficiency complementation group 1 (includes overlapping antisense sequence)
aldo-keto reductase family 7, member A2 (aflatoxin aldehyde reductase)
serine hydroxymethyltransferase 2 (mitochondrial)
putative nucleic acid binding protein RY-1
RAS protein activator like 1 (GAP1 like)
phosphorylase, glycogen; brain
ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)
Ran GTPase activating protein 1
amino-terminal enhancer of split
calpain, small subunit 1
gb:NM_005703.2 /DEF=Homo sapiens upstream regulatory element binding protein 1 (UREB1), mRNA, /FEA=CDS /GEN=UREB1 /PROD=upstream regulatory element bindin
gb:AF092132.1 /DEF=Homo sapiens PAK2 mRNA, complete cds, /FEA=mRNA /PROD=PAK2 /DB_XREF=gi:5138913 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete
solute carrier family 7, (cationic amino acid transporter, y+ system) member 11
hypothetical protein
integrin-linked kinase-associated serine/threonine phosphatase 2C
phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminimidazole synthetase
ADP-ribosyltransferase (NAD+; poly(ADP-ribose) polymerase)-like 2
v-rat murine sarcoma 3811 viral oncogene homolog 1
SH3-domain, GRB2-like, endophilin B1
GRB2-associated binding protein 1
F-box only protein 9
ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast)
RNA processing factor 1
proteasome (prosome, macropain) 26S subunit, ATPase, 5
transcription factor 2, hepatic; LF-83; variant hepatic nuclear factor
microtubule-associated protein 4
CD2 antigen (cytoplasmic tail) binding protein 2
antigen identified by monoclonal antibody Ki-67
hypothetical protein FLJ10241
casein kinase 2, alpha 1 polypeptide
HCR (e-helix coiled-coil rod homologue)
hypothetical protein FLJ23591
ATP-binding cassette, sub-family B (MDR/TAP), member 7
hypothetical protein FLJ12529
breast cancer 1, early onset
HIV-1 Tat interactive protein 2, 30 kDa
high density lipoprotein binding protein (vigin)
hypothetical protein MGC14258
opoid growth factor receptor
chemokine-like factor 1
stromal antigen 2
actin A receptor, type IIB
hypothetical protein FLJ12443
glycogen synthase kinase 3 alpha
CCR4-NOT transcription complex, subunit 3
myozenin
cyclic AMP phosphoprotein, 19 kD
synapsin II
nucleosome assembly protein 1-like 1
zinc finger protein 91 (HPP7, HTF10)
arsenate resistance protein ARS2
ornithine decarboxylase antizyme 1
hypothetical protein FLJ10842
nuclear respiratory factor 1
Consensus includes gb:AK000345.1 /DEF=Homo sapiens cDNA FLJ20338 fts, clone HEP12179, /FEA=mRNA /DB_XREF=gi:7020368 /UG=Hs.152677 Homo sapiens cDNA FI
PAL-1 mRNA-binding protein
putative c-Myc-responsive
hypothetical protein DKFZp566F2423
Consensus includes gb:AK026008.1 /DEF=Homo sapiens cDNA: FLJ22355 fts, clone HRC06344, /FEA=mRNA /DB_XREF=gi:10438707 /UG=Hs.288283 Homo sapiens cDNA
v-myc myelocytomatosis viral oncogene homolog (avian)
syntaxin 7
casein kinase 1, gamma 3

BC001288 1.46
NM_001983 1.46
NM_003689 1.46
AW190316 1.46
BG252325 1.46
NM_004658 1.46
NM_002862 1.46
AA602532 1.46
BE378408 1.46
NM_001130 1.46
NM_001749 1.46
NM_005703 1.46
AF092132 1.46
AB040875 1.46
NM_016458 1.46
AY024365 1.46
BE966876 1.46
NM_005484 1.46
NM_001654 1.46
AF257318 1.46
NM_002039 1.46
AI954041 1.46
NM_003348 1.46
NM_025065 1.46
AF055309 1.46
NM_000458 1.46
NM_002375 1.46
NM_006110 1.46
AU132185 1.46
AF768100 1.46
AI631874 1.46
AB029343 1.46
NM_025076 1.46
AB005289 1.46
NM_024811 1.46
AF005068 1.46
NM_006410 1.46
NM_005336 1.45
U79256 1.45
NM_007346 1.45
NM_016326 1.45
NM_006603 1.45
NM_001106 1.45
NM_024830 1.45
NM_019684 1.45
AF180474 1.45
NM_021245 1.45
BC003418 1.45
AW139618 1.45
AI888672 1.45
NM_003430 1.45
NM_015908 1.45
D87914 1.45
AJ278150 1.45
NM_005011 1.45
AF151813 1.45
AF040105 1.45
BE975139 1.45
AK026008 1.45
NM_002467 1.45
NM_003569 1.45
NM_004384 1.45

huntingtin interacting protein 1 1.45
 hypothetical protein FLJ12700 1.45
 succinate-CoA ligase, GDP-forming, beta subunit 1.45
 ADP-ribosylation factor-like 2 1.45
 replication protein A2 (32kD) 1.45
 phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole synthetase 1.45
 platelet-activating factor acetylhydrolase, isoform 1b, alpha subunit (45kD) 1.45
 hypothetical protein MGC2722 1.45
 cachectin, EGF LAG seven-pass G-type receptor 3 (flamingo homolog, Drosophila) 1.45
 KIAA0637 gene product 1.45
 Consensus includes gb:AW371924 /FEA=EST /DB_XREF=gi:6876578 /DB_XREF=est:RC4-BT0311-251199-012-a07 /UG=Hs.313848 ESTs, Moderately similar to MGN_HUM_AW371924 1.45
 CGI-99 protein 1.45
 poly(pyrimidine) tract binding protein (heterogeneous nuclear ribonucleoprotein 1) 1.45
 bedlin 1 (coiled-coil, myosin-like BCL2 interacting protein) 1.45
 endoplasmic reticulum chaperone SIL1, homolog of yeast 1.44
 homolog of mouse quaking QKI (KH domain RNA binding protein) 1.44
 propionyl Coenzyme A carboxylase, beta polypeptide 1.44
 FUS interacting protein (serine-arginine rich) 1 1.44
 hypothetical protein FLJ21952 1.44
 hypothetical protein FLJ20374 1.44
 polymerase (RNA) II (DNA directed) polypeptide E (25kD) 1.44
 interleukin 10 receptor, beta 1.44
 COP9 subunit 6 (MOV34 homolog, 34 kD) 1.44
 protein tyrosine phosphatase, non-receptor type 2 1.44
 Cluster Ind. AA160496:73a03.a1 Homo sapiens cDNA, 3 end /clone=IMAGE:592492 /clone_end=3 /gb=AA160498 /ug=Hs.110407 /len=542 1.44
 mitofusin 1 1.44
 splicing factor, arginine/serine-rich 5 1.44
 defensin, beta 2 1.44
 GM2 ganglioside activator protein 1.44
 general transcription factor IIIC, polypeptide 5 (63kD) 1.44
 VRK3 for vaccinia related kinase 3 1.44
 Cluster Ind. A1144299:qb59n08.x1 Homo sapiens cDNA, 3 end /clone=IMAGE:1704443 /clone_end=3 /gb=A1144299 /ug=Hs.106843 /len=765 1.44
 casein kinase 1, alpha 1 1.44
 Kruppel-like factor 12 1.44
 DKF ZP434D156 protein 1.44
 phosphatidylinositol transfer protein, beta 1.44
 thoredoxin, mitochondrial 1.44
 host cell factor C1 (VP16-accessory protein) 1.44
 casein, kappa 1.44
 rat guanine nucleotide dissociation stimulator 1.44
 gb:NM_015930.1 /DEF=Homo sapiens vanilloid receptor-like protein 1 (VRL-1), mRNA. /FEA=mRNA /GEN=VRL-1 /PROD=vanilloid receptor-like protein 1 /DB_XREF=gi:77067 /NM_015930 1.44
 scaffold attachment factor 8 1.44
 myelin protein zero-like 1 1.44
 hypothetical protein FLJ10546 1.44
 phosphatidylinositol 4-kinase, catalytic, alpha polypeptide 1.44
 Ndc66 (D. melanogaster)-like protein 1.44
 KIAA1253 protein 1.44
 regulator of nonsense transcripts 1 1.44
 Consensus includes gb:AL133053.1 /DEF=Homo sapiens mRNA; cDNA DKF ZP434H1215 (from clone DKF ZP434H1215); partial cds. /FEA=mRNA /GEN=DKF ZP434H1215 /P AL133053 1.44
 Consensus includes gb:BF056534 /FEA=EST /DB_XREF=gi:10810430 /DB_XREF=est:7406609.x1 /CLONE=IMAGE:34413441 /UG=Hs.123469 ESTs, Weakly similar to AF208 /BF056534 1.44
 BCL2 adenovirus E1B 19kD interacting protein 1 1.44
 Consensus includes gb:BE680028 /FEA=EST /DB_XREF=gi:10329604 /DB_XREF=est:60149326 /F1 /CLONE=IMAGE:3895407 /UG=Hs.84775 Human transposon-like element BE680028 1.44
 Consensus includes gb:AW081820 /FEA=EST /DB_XREF=gi:6036972 /DB_XREF=est:xb55d12.x1 /CLONE=IMAGE:2580215 /UG=Hs.287317 ESTs, Weakly similar to AS9252 /AW081820 1.44
 poly(A) binding protein, cytoplasmic 4 (inducible form) 1.44
 polymerase (DNA directed), epsilon 1.44
 survival of motor neuron protein interacting protein 1 1.44
 DKF Zp556F1019 protein 1.44
 golgi associated, gamma adaptin ear containing, ARF binding protein 2 1.44
 major histocompatibility complex, class II, DM beta 1.44
 capping protein (actin filament) muscle Z-line, beta 1.44
 Consensus includes gb:AU145293 /FEA=EST /DB_XREF=gi:11006814 /DB_XREF=est:AU145293 /CLONE=HEMBA1004405 /UG=Hs.61773 Homo sapiens cDNA FLJ11648 fl AU145293 1.44
 vesicle-associated membrane protein 2 (synaptobrevin 2) 1.44
 gb:NM_020382.1 /DEF=Homo sapiens PRSET domain containing protein 07 (SET07), mRNA. /FEA=mRNA /GEN=SET07 /PROD=PRSET domain containing protein 07 /DB_X NM_020382 1.44

KIAA0540 protein	AB011112	1.44
ribosomal protein S23	NM_001025	1.44
KIAA0816 protein	AB014516	1.44
thioredoxin	AF313911	1.44
polymerase (RNA) II (DNA directed) polypeptide C (33kD)	BC000409	1.44
signal transducer and activator of transcription 3 (acute-phase response factor)	BC000627	1.44
similar to H2A histone family, member A (H. sapiens)	BC001193	1.43
hypothetical protein FLJ13660 similar to CDK5 activator-binding protein C53	NM_025197	1.43
presenilin associated rhomboid-like protein	NM_018622	1.43
glucocorticoid receptor DNA binding factor 1	AU159484	1.43
guanine nucleotide binding protein (G protein), alpha activating activity polypeptide O	NM_020988	1.43
a disintegrin and metalloproteinase domain 10	NM_001110	1.43
thymidine kinase 1, soluble	NM_003258	1.43
programmed cell death 8 interacting protein	NM_013374	1.43
ubiquinol-cytochrome c reductase (8.4kD) subunit	NM_006830	1.43
zinc finger protein 313	NM_018683	1.43
chromobox homolog 4 (Pc class homolog, Drosophila)	NM_003655	1.43
Consensus includes gp:AV728528 /FEA=EST/DB_XREF=gi:10837947 /DB_XREF=est:AV728528 /CLONE=HTCBBDD04 /UG=Hs.62264 KIAA0937 protein	AV728526	1.43
KIAA0179 protein	AI400490	1.43
RAD1 homolog (S. pombe)	D79998	1.43
proliferating cell nuclear antigen	AF074717	1.43
NADH dehydrogenase (ubiquinone) Fe-S protein 2 (49kD) (NADH-coenzyme Q reductase)	NM_002592	1.43
mitogen-activated protein kinase 1	NM_004550	1.43
katanin p80 (WD40-containing) subunit B 1	NM_002745	1.43
MAGEF1 protein	NM_005886	1.43
Jun D proto-oncogene	NM_022149	1.43
nurt-related transcription factor 3	AI339541	1.43
CHK1 checkpoint homolog (S. pombe)	NM_004350	1.43
synapsin 10	NM_001274	1.43
solute carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc) transporter), member 3	NM_003765	1.43
hypothetical protein FLJ11808	NM_012243	1.43
phosphoglycerate kinase 1	NM_024824	1.43
polyamine-modulated factor 1	NM_000291	1.43
endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 7	NM_007221	1.43
N-myristoyltransferase 1	NM_012152	1.43
Consensus includes gp:AW575379 /FEA=EST/DB_XREF=gi:7246918 /DB_XREF=est:U1-HF-8L0-acc-e-05-O-U1.s1 /CLONE=IMAGE:3060969 /UG=Hs.6551 ATPase, H ⁺ trans AW575379	AF020500	1.43
O-linked N-acetylglucosamine (GlcNAc) transferase (UDP-N-acetylglucosamine polypeptide-N-acetylglucosaminyl transferase)	BF001665	1.43
glucosaminyl (N-acetyl) transferase 2, branching enzyme	L19659	1.43
v-abl Abelson murine leukemia viral oncogene homolog 1	NM_005157	1.43
IKK-related kinase epsilon; inducible IkappaB kinase	NM_014002	1.43
methionine-tRNA synthetase	AA621558	1.43
RET-silencing transcription factor	NM_005612	1.43
ubiquitin-conjugating enzyme HBUCE1	NM_015983	1.43
sulfotransferase family, cytosolic, 1C, member 2	NM_006588	1.43
adenylyate kinase 2	AL513611	1.43
Consensus includes gp:AI658004 /FEA=EST/DB_XREF=gi:5511620 /DB_XREF=est:wj69009.x1 /CLONE=IMAGE:2408057 /UG=Hs.284277 Homo sapiens immunoglobulin m	AI658004	1.43
hypothetical protein FLJ21986	NM_024913	1.43
reticulocalbin 2, EF-hand calcium binding domain	BC004892	1.43
yeast Sec31p homolog	AB020712	1.43
cholinergic receptor, nicotinic, alpha polypeptide 4	X89745	1.43
LGN protein	NM_013296	1.43
hypothetical protein MG20727	N32831	1.43
death effector domain-containing	AF064605	1.43
polypyrimidine tract binding protein (heterogeneous nuclear ribonucleoprotein l)	NM_002819	1.43
fragile X mental retardation, autosomal homolog 1	NM_005087	1.43
solute carrier family 25 (carnitine/acylcarnitine translocase), member 20	BC001689	1.43
meningioma expressed antigen 6 (coiled-coil proline-rich)	U73682	1.43
hypothetical protein FLJ21062	NM_024788	1.43
glutaminase	AB020645	1.43
RNA binding motif protein 9	AW149379	1.43
stress-associated endoplasmic reticulum protein 1; ribosome associated membrane protein 4	AL136807	1.43
SET domain and mariner transposase fusion (gmm)	NM_006515	1.42

epoxide hydrolase 2, cytoplasmic	AF233336	1.42
KIAA0683 gene product	AK022669	1.42
protein inhibitor of activated STAT protein PIASy	NM_015897	1.42
KIAA0429 gene product	AF116674	1.42
G protein pathway suppressor 2	AL157493	1.42
hypothetical protein FLJ21736	NM_024922	1.42
polycystic kidney disease 2-like 1	NM_016112	1.42
cholinergic receptor, nicotinic, alpha polypeptide 7	U62436	1.42
leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2	U62278	1.42
Consensus includes gb:R37104 /FEA=EST /DB_XREF=gi:794560 /DB_XREF=est:53e12.a1 /CLONE=IMAGE:25731 /UG=Hs.13417 Homo sapiens clone 24670 mRNA seq	R37104	1.42
insulin-degrading enzyme	N22903	1.42
killer cell lectin-like receptor subfamily A, member 1	NM_006611	1.42
serum response factor (c-fos serum response element-binding transcription factor)	NM_003131	1.42
calretin	U64869	1.42
ribosomal protein L41	NM_021104	1.42
succinate-CoA ligase, GDP-forming, alpha subunit	NM_003849	1.42
ubiquitin-like factor E4B (UFD2 homolog, yeast)	AW241715	1.42
hypothetical protein	AA748492	1.42
phosphorylase kinase, gamma 2 (testis)	NM_000294	1.42
deoxythymidylate kinase (thymidylate kinase)	NM_012145	1.42
cell death regulator even	NM_020371	1.42
hypothetical protein FLJ10547	AL049795	1.42
endonuclease G-like 2	NM_005728	1.42
Cluster incl. AA32670:nl39a05.a1 Homo sapiens cDNA, 3' end /clone=IMAGE:979184 /clone_end=3' /gb=AA522670 /gi=2263382 /ug=Hs.7247 /len=684	AA522670	1.42
prolylcarboxypeptidase (angiotensinase C)	NM_005040	1.42
transducin-like enhancer of split 4 (E(spl) homolog, Drosophila)	AL358975	1.42
H2A histone family, member P	NM_021064	1.42
Consensus includes gb:AJ275383 /DEF=Homo sapiens partial IGVH3 gene for immunoglobulin heavy chain V region, case 1, cell Mo IV 72 /FEA=CDS /DB_XREF=gi:7573027	AJ275383	1.42
immunoglobulin lambda variable 3-10	D87021	1.42
caspase 6, apoptosis-related cysteine protease	BC000305	1.42
COI-26 protein	NM_015954	1.42
ubiquitin-conjugating enzyme E2E 3 (UBC4/5 homolog, yeast)	AB017644	1.42
translocase of inner mitochondrial membrane 23 homolog (yeast)	NM_006327	1.42
beta-amyloid binding protein precursor	AF353990	1.42
calreticulin	AD000092	1.42
Sec3-like	AK023461	1.42
cell division cycle 2-like 1 (PITSLRE proteins)	NM_001787	1.42
peroxisomal biogenesis factor 6	D83703	1.42
hypothetical protein BK1048E9.5	AL041451	1.42
testis specific protein, Y-linked	M94893	1.42
KIAA0876 protein	AK026096	1.42
heat shock 60kD protein 1 (chaperonin)	NM_002156	1.42
succinate-CoA ligase, GDP-forming, beta subunit	AF131748	1.42
IMP (inosine monophosphate) dehydrogenase 2	NM_000884	1.42
26S proteasome-associated pcd1 homolog	NM_005805	1.42
ATP-binding cassette, sub-family C (CFTR/MRP), member 8	AF087138	1.42
DEK oncogene (DNA binding)	NM_003472	1.42
KIAA0184 protein	D80006	1.42
nuclear pore complex interacting protein	NM_006985	1.42
dihydropyrimidinase-like 4	AW090187	1.42
cyclin G associated kinase	NM_005255	1.42
nucleoporin-like protein 1	NM_007342	1.42
glutaminase	NM_014905	1.42
intercellular adhesion molecule 1 (CD54), human rhinovirus receptor	NM_000201	1.42
hypothetical protein MGC3077	AK021779	1.42
Consensus includes gb:BG170659 /FEA=EST /DB_XREF=gi:12677362 /DB_XREF=est:60232353.1 /CLONE=IMAGE:4426926 /UG=Hs.226138 Homo sapiens mRNA, cDNA BG170659	BG170659	1.42
hypothetical protein FLJ20850	AJ74084	1.42
truncobiose repeat containing 12	BE860591	1.41
actin, alpha, cardiac muscle	NM_005159	1.41
TNF-induced protein	NM_014350	1.41
mitogen-activated protein kinase-activated protein kinase 3	NM_004635	1.41
hypothetical protein FLJ20384	AF268167	1.41
putative methyltransferase	BC001650	1.41

gb|U18810.1/DEF=Human PACAP type-3VIP type-2 receptor mRNA, complete cds. /FEA=mRNA /PROD=PACAP type-3VIP type-2 receptor /DB_XREF=gi:1884934 /FL=gb|U: U18810 1.41
 potassium voltage-gated channel, Isk-related family, member 1-like 1.41
 proteasome (prosome, macropain) activator subunit 3 (PA28 gamma: Ki) 1.41
 hypothetical protein FLJ20607 1.41
 ribosomal protein S28 1.41
 homolog of mouse quaking OK1 (KH domain RNA binding protein) 1.41
 Consensus includes gb|AA0203487 /FEA=EST /DB_XREF=gi:1799460 /DB_XREF=est:zx53403.1 /CLONE=IMAGE:446213 /UG=Hs.314363 ESTs 1.41
 putative G-protein coupled receptor 1.41
 H3 histone, family 3B (H3.3B) 1.41
 KIAA0870 protein 1.41
 MAGIE-E1 protein 1.41
 integral membrane protein 2B 1.41
 hypothetical protein FLJ2127 1.41
 FK506 binding protein 12-repamycin associated protein 1 1.41
 PTD008 protein 1.41
 hypothetical protein FLJ20850 1.41
 transglutaminase 1 (K polypeptide epidermal type 1, protein-glutamine-gamma-glutamyltransferase) 1.41
 mentholase (diphospho) decarboxylase 1.41
 N-acetylphosphatase endohydrolase (acid ceramidase)-like 1.41
 cysteine binding protein 1.41
 Consensus includes gb|AJ02559 /DEF=Homo sapiens 6M1-4P*01 pseudogene, cell line LG2 /FEA=CDS /DB_XREF=gi:12054348 /UG=Hs.307110 Homo sapiens 6M1-4P*01 1.41
 H2B histone family, member F 1.41
 golgi associated, gamma adaptin ear containing, ARF binding protein 2 1.41
 mitofusin 2 1.41
 HP1-BP74 1.41
 KIAA1888 protein 1.41
 ADP-ribosylation factor-like 6 interacting protein 1.41
 ADP-ribosylation factor-like 1 1.41
 arylsulatase E (chondrocyte-specific punctate 1) 1.41
 vesicle docking protein p115 1.41
 gb|BC001121.1/DEF=Homo sapiens. Similar to deoxyguanosine kinase /DB_XREF=gi:12054348 /UG=Hs.307110 Homo sapiens 6M1-4P*01 1.41
 hypothetical protein FLJ23309 1.41
 proteasome (prosome, macropain) subunit, beta type, 4 1.41
 mitofusin 2 1.41
 Rab geranylgeranyltransferase, alpha subunit 1.41
 Consensus includes gb|AU144887 /FEA=EST /DB_XREF=gi:11006408 /DB_XREF=est:AU144887 /CLONE=HEMBA1003278 /UG=Hs.301800 Homo sapiens cDNA FLJ11568 1.41
 H-appea-B-interacting Ras-like protein 2 1.41
 homeo box D3 1.41
 DEAH1 (Asp-Glu-Ala-AspHis) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophycin) 1.41
 autoantigen 1.41
 hypothetical protein MGC2840 similar to a putative glucosyltransferase 1.41
 GLI-Kruppel family member HKR3 1.41
 hypothetical protein MGC2780 1.40
 tubulin-specific chaperone d 1.40
 KIAA0848 protein 1.40
 tropomodulin 3 (ubiquitous) 1.40
 hypothetical protein FLJ10312 1.40
 APO12 autophagy 12-like (S. cerevisiae) 1.40
 RNA binding motif, single stranded interacting protein 1 1.40
 hypothetical protein FLJ21213 1.40
 interferon, alpha 17 1.40
 sekute carrier family 16 (monocarboxylic acid transporters), member 3 1.40
 ReoQ protein-like 5 1.40
 HOM-TES-45 tumor antigen 1.40
 keratophen (imporin) beta 3 1.40
 Consensus includes gb|BC0285011 /FEA=EST /DB_XREF=gi:13035541 /DB_XREF=est:60209278F1 /CLONE=IMAGE:4538275 /UG=Hs.12702 Homo sapiens mRNA: cDNA t BC0285011 1.40
 CGI-49 protein 1.40
 triple functional domain (PTPRF interacting) 1.40
 Homer, neuronal immediate early gene, 3 1.40
 tropinin C, slow 1.40
 erlapin 1 1.40
 activin A receptor, type IB 1.40

major histocompatibility complex, class I, J (pseudogene)	M80469	1.40
cytochrome P450, subfamily 1J (arachidonic acid epooxygenase) polypeptide 2	NM_000775	1.40
bilirubin reductase B (flavin reductase (NADPH))	NM_000713	1.40
chromosome 21 open reading frame 66	NM_016631	1.40
hypothetical protein FLJ14888	AF161382	1.40
entylin repeat and SOCS box-containing 1	AF055024	1.40
hypothetical protein FLJ10581	NM_016146	1.40
colony stimulating factor 1 (macrophage)	NM_000757	1.40
hydroxyacyl-Coenzyme A dehydrogenase/3-ketocyl-Coenzyme A thiolase/acyl-Coenzyme A hydratase (trifunctional protein), beta subunit	NM_000183	1.40
IQ motif containing GTPase activating protein 1	D29640	1.40
zinc finger protein 95 homolog (mouse)	BF196831	1.40
emil1 sequence (mammary tumor and squamous cell carcinoma-associated (p80/85 src substrate))	BG475299	1.40
CASP8 and FADD-like apoptosis regulator	AF015451	1.40
YME1-like 1 (S. cerevisiae)	AF070656	1.40
elcC homolog 2 (E. coli)	NM_018127	1.40
RAN binding protein 18	NM_015024	1.40
LRAP for lysophosphatidic acid phosphatase	NM_016361	1.40
hypothetical protein DKFp762M115	BC024777	1.40
calumenin	BF936365	1.40
ret finger protein	NM_006510	1.40
dishevelly/peptidase III	NM_005700	1.40
plakophilin 4	NM_003628	1.40
acyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase)	AW072302	1.40
ribosomal protein L37a	NM_000988	1.40
intercellular adhesion molecule 1 (CD54), human rhinovirus receptor	AI608725	1.40
splicing factor 1	NM_004630	1.40
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump), beta polypeptide, 56/58kD, isoform 1 (Renal tubular acidosis with deafness)	NM_001692	1.40
cell division cycle 23C	NM_001790	1.40
novel putative protein similar to YIL091C yeast hypothetical 84 kD protein from SGA1-KTR7	N30910	1.40
CD64 antigen (leukocyte antigen)	NM_003874	1.40
polymerase (DNA-directed), alpha (70kD)	NM_002689	1.39
RNA polymerase I subunit	NM_004875	1.39
SET binding factor 1	U93181	1.39
receptor (TNFRSF)-interacting serine-threonine kinase 1	U50062	1.39
RAD23 homolog A (S. cerevisiae)	NM_005053	1.39
splicing factor 3b, subunit 1, 155kD	NM_012433	1.39
phosphatase and leucine homolog (mutated in multiple advanced cancers 1), pseudogene 1	NM_018346	1.39
hypothetical protein FLJ11164	AF023139	1.39
hypothetical protein FLJ10592	NM_022914	1.39
hypothetical protein 24638	BF439241	1.39
ab1-interactor 12 (SH3-containing protein)	NM_018147	1.39
proteasome (prosome, macropain) 26S subunit, non-ATPase, 12	BC002840	1.39
LJM protein (similar to rat protein kinase C-binding enigma)	AF260261	1.39
vacuolar protein sorting 18 (yeast)	AI446530	1.39
Consensus includes gb-M74089.1 (DEF-Human TB1 gene mRNA, 3 end. /FEA-mRNA /GEN=TB1 /DB_XREF=gi:182400 /UG=Hs.75639 Human TB1 gene mRNA, 3 end	NM_008457	1.39
CCR4-NOT transcription complex, subunit 2	NM_022575	1.39
primase, polypeptide 1 (49kD)	M74089	1.39
PCTAIRE protein kinase 2	BG105204	1.39
vesicular endonuclease growth factor	NM_000946	1.39
ceroid-lipofuscinosis, neuronal 5	NM_002395	1.39
G protein-coupled receptor 38	M27281	1.39
a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme)	NM_006483	1.39
tetrapsan 3	NM_001507	1.39
TAF9 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 32 kD	AI797833	1.39
H1 histone family, member 0	BC000704	1.39
KIAA0020 gene product	NM_003187	1.39
SH3 domain binding glutamic acid-rich protein like	Z97630	1.39
hypothetical protein FLJ10120	NM_014878	1.39
hypothetical protein FLJ23022	AL51518	1.39
Cluster Ind. AW001777:wd4f09.x1 Homo sapiens cDNA, 3' end (clone=IMAGE-2498233 /clone_end=3' (gb=AW001777 /gi=5848693 /ug=Hs.233841 /len=465	NM_018001	1.39
KIAA0019 protein	NM_025051	1.39
	AW001777	1.39
	AB020626	1.39

protein kinase, AMP-activated, beta 1 non-catalytic subunit
KIAA0456 protein
zinc finger protein 117 (HPF9)
RAP2A, member of RAS oncogene family
peptide-N-glycanase similar to yeast PNG1
sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase)
KIAA0372 protein
Consensus includes gb:BG337255 /FEA=EST /DB_XREF=gi:13529117 /DB_XREF=est:602565318F1 /CLONE=IMAGE:4689748 /UG=Hs.82646 DnaJ (Hsp40) homolog, subfamily A (Asp-Glu-Ala-Asp/His) box polypeptide 3
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 3
general transcription factor II, 1
neuronal PAS domain protein 2
DNFZP568F1916 protein
30 kDa protein
KIAA0076 gene product
uroporphyrinogen III synthase (congenital erythropoietic porphyria)
heparan sulfate 6-O-sulfotransferase
RNA binding motif, single stranded interacting protein 1
signal recognition particle 9kD
Consensus includes gb:AK023154.1 /DEF=Homo sapiens cDNA FLJ13092 (is. clone NT2RP002147) /FEA=mRNA /DB_XREF=gi:10434948 /UG=Hs.172035 hypothetical protein AK023154
nuclear cap binding protein subunit 2, 20kD
LPS-induced TNF-alpha factor
filamin B, beta (actin binding protein 278)
myosin, light polypeptide 4, alkali; atrial; embryonic
APG3 autophagy 5-like (S. cerevisiae)
adducin 1 (alpha)
mitochondrial ribosomal protein L15
hypothetical protein FLJ21080
splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila)
reqlum, apoptotic response zinc finger gene
desmin
F-box only protein 9
tyrosyl-tRNA synthetase
KIAA0365 gene product
chromosome 1 open reading frame 28
JAK binding protein
B-cell CLL/lymphoma 7A
hypothetical protein MGC3180
sorting nexin 6
hypothetical protein MGC10882
lamin B2
inositol polyphosphate 4-phosphatase, type I, 107kD
nuclear pore complex interacting protein
chromosome 5 open reading frame 4
cleavage and polyadenylation specific factor 4, 30kD subunit
autotransferase family, cytosolic, 1A, phenol-preferring, member 3
translocation protein 1
ribokinase
Consensus includes gb:AL008627 /DEF=Human DNA sequence from PAC 130G2 on chromosome 6p22.2-22.3. Contains ribosomal protein L29 pseudogene, ESTs and STSs
ribonuclease P (14kD)
metastasis associated 1
protein phosphatase 1, regulatory (inhibitor) subunit 2
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophycin)
hypothetical protein FLJ2427
tyrosylprotein sulfotransferase 2
foditin 1
anaphase-promoting complex subunit 10
proline oxidase homolog
transmembrane 7 superfamily member 2
carboxypeptidase D
DNFZP34N043 protein
chondrohypoxanthine 81
eukaryotic translation initiation factor 3, subunit 7 (zeta, 66/67kD)
hypothetical protein FLJ20186

BC001007 1.38
AA742261 1.38
NM_024498 1.38
AI02106 1.38
NM_018297 1.38
AI743792 1.38
AI629033 1.38
BG537255 1.38
AF061327 1.38
AC004863 1.38
NM_002518 1.38
BE645501 1.38
NM_018447 1.38
NM_014780 1.38
NM_000375 1.38
NM_004807 1.38
NM_016637 1.38
NM_003133 1.38
AK023154 1.38
BC001255 1.38
NM_004862 1.38
AV12733 1.38
X58851 1.38
AK001889 1.38
BE688639 1.38
NM_014175 1.38
NM_022743 1.38
AI871457 1.38
NM_006268 1.38
NM_001927 1.38
AF176704 1.38
W27419 1.38
AV738039 1.38
AW192876 1.38
AI056051 1.38
NM_020993 1.38
NM_024041 1.38
NM_021249 1.38
BC004952 1.38
M94363 1.38
NM_004027 1.38
AA308653 1.38
H93077 1.38
NM_006693 1.38
U08032 1.38
U83239 1.38
NM_022128 1.38
AL008627 1.38
AV242755 1.38
BE561596 1.38
BF966540 1.38
NM_001357 1.38
BF129339 1.38
NM_003595 1.38
AF085357 1.38
NM_014885 1.38
AA074145 1.38
AF086304 1.38
NM_001304 1.38
BG538800 1.38
AW584011 1.38
NM_003753 1.38
NM_017703 1.38

ubiquitin-conjugating enzyme E2D 3 (UBC4/5 homolog, yeast) 1.38
 chromosome 8 open reading frame 7 1.38
 nuclear pore complex interacting protein 1.38
 KIAA1083 protein 1.38
 hypothetical protein MGC2454 1.38
 SRp25 nuclear protein 1.38
 proline rich 1 1.38
 nucleolar cytosine-rich protein 1.38
 mitogen-activated protein kinase 1 1.38
 MCH43, mitochondrial maintenance deficient 3 (S. cerevisiae) 1.38
 hypothetical protein FLJ21877 1.38
 Bruton agammaglobulinemia tyrosine kinase 1.38
 RNA polymerase I transcription factor RRN3 1.38
 origin recognition complex, subunit 2-like (yeast) 1.38
 Consensus includes gb:AV700332 /FEA=EST /DB_XREF=gi:10302303 /DB_XREF=gi:10302303 /CLONE=GKCDIF.11 /UG=Hs.243501 Homo sapiens cDNA FLJ20738 fls, ck AV700332 1.38
 butyrate-induced transcript 1 1.38
 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASH1) 1.38
 CD3D antigen, delta polypeptide (T173 complex) 1.38
 protein tyrosine phosphatase, non-receptor type 6 1.38
 proenkephalin 1.38
 retinol x receptor interacting protein 1.38
 solute carrier family 35 (UDP-galactose transporter), member 2 1.38
 hypothetical protein FLJ20294 1.38
 KIAA0592 protein 1.38
 Consensus includes gb:AL080233.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586L111 (from clone DKFZp586L111) /FEA=mRNA /DB_XREF=gi:5262726 /UG=Hs.326580 H AL080233 1.38
 transcriptional adaptor 2 (ADA2 homolog, yeast)-like 1.38
 apoptosis-associated spect-like protein containing a CARD 1.38
 Bicaudal D homolog 1 (Drosophila) 1.38
 SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome) 1.37
 pre-B lymphocyte gene 3 1.37
 hypothetical protein FLJ10509 1.37
 nuclear VCP-like 1.37
 DC13 protein 1.37
 zinc finger RNA binding protein 1.37
 membrane metallo-endopeptidase (neutral endopeptidase, enkephalinase, CALLA, CD10) 1.37
 activated RNA polymerase II transcription colactor 4 1.37
 HLA class II region expressed gene KE2 1.37
 karyopherin (importin) beta 1 1.37
 histone acetyltransferase 1.37
 poly(A) binding protein, cytoplasmic, pseudogene 3 1.37
 Consensus includes gb:T33068 /FEA=EST /DB_XREF=gi:615166 /DB_XREF=est:EST56576 /UG=Hs.7101 anaphase-promoting complex subunit 5 1.37
 hypothetical protein FLJ14007 1.37
 hypothetical protein FLJ10878 1.37
 nuclear pore complex interacting protein 1.37
 cornichon-like 1.37
 gb:NM_001514.2 /DEF=Homo sapiens general transcription factor IIB (GTF2B), mRNA /FEA=mRNA (GEN=GTF2B) /PROD=general transcription factor IIB /DB_XREF=gi:1343 NM_001514 1.37
 Consensus includes gb:BG255188 /FEA=EST /DB_XREF=gi:12765004 /DB_XREF=est:802369727F1 /CLONE=IMAGE:4477807 /UG=Hs.3343 phosphoglycerate dehydrogenase BG255188 1.37
 mitochondrial ribosomal protein S30 1.37
 hypothetical protein FLJ20258 1.37
 replication protein A1 (70kD) 1.37
 glucocorticoid modulatory element binding protein 1 1.37
 transmembrane 4 superfamily member 7 1.37
 trophinin associated protein (tastin) 1.37
 cytosolic acyl coenzyme A thioester hydrolase 1.37
 KIAA0652 gene product 1.37
 NG23 protein 1.37
 neuroblastoma binding protein 8 1.37
 RAB28, member RAS oncogene family 1.37
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 1.37
 GS2 gene 1.37
 uncharacterized bone marrow protein BM045 1.37
 MLN51 protein 1.37
 acyl-Coenzyme A acyltransferase 2 (mitochondrial 3-oxoacyl-Coenzyme A thiolase) 1.37

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forward box O1A (mabdomyosarcoma)
 serine palmitoyltransferase, long chain base subunit 2
 RAB35, member RAS oncogene family
 NS1-associated protein 1
 inhibitor of DNA binding 2, dominant negative helix-loop-helix protein
 DNA segment, single copy, probe pH4 (transforming sequence, thyroid-1,
 general transcription factor IIIA)
 carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydroorotase
 phosphoprotein enriched in astrocytes 15
 exosome component Rnp41
 inositol 1,4,5-trisphosphate receptor, type 3
 dual specificity phosphatase 1
 hypothetical protein FLJ11338
 methyl-CpG binding domain protein 3
 hypothetical protein FLJ22494
 hydromethylbilane synthase
 hypothetical protein R33729_1
 phosphoinositide-3-kinase, regulatory subunit, polypeptide 1 (p85 alpha)
 calcium channel, voltage-dependent, P/Q type, alpha 1A subunit
 steroidogenic acute regulatory protein related
 dyenin, cytoplasmic, light polypeptide
 protein phosphatase 2, regulatory subunit B (B56), gamma isoform
 TGF-beta-induced anti-apoptotic factor 1
 KIAA0365 gene product
 hypothetical protein FLJ21459
 acetyl-Coenzyme A transporter
 hypothetical protein FLJ10652
 bone morphogenetic protein 15
 leucine zipper transcription factor-like 1
 KIAA0683 gene product
 KIAA0676 protein
 Consensus includes gb:AI007083 /FEA=EST /DB_XREF=gi:6497611 /DB_XREF=est:PM-BT134-050499-650 /UG=Hs.124620 ESTs
 Consensus includes gb:AL133015.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434O2417 (from clone DKFZp434O2417); partial cds. /FEA=mRNA /GEN=DKFZp434O2417 IF AL133015
 amplified in osteosarcoma
 KIAA0794 protein
 similar to vaccinia virus HindIII K4L ORF
 methyl-CpG binding domain protein 1
 hypothetical protein FLJ20560
 splicing factor, arginine/serine-rich 6
 Consensus includes gb:AJ295618 /DEF=Homo sapiens FISH gene for putative ATPases, exons 1 and 2 and join CDS /FEA=mRNA /DB_XREF=gi:8506352 /UG=Hs.206521 Y AJ295618
 eriodine homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila)
 zinc finger protein, subfamily 1A, 1 (Ikroos)
 gastrin-releasing peptide receptor
 solute carrier family 5 (sodium/glucose cotransporter), member 2
 hypothetical protein FLJ10335
 protein kinase, cAMP-dependent, regulatory, type II, beta
 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
 mitochondrial ribosomal protein S18A
 lysosomal-associated membrane protein 1
 SRY (sex determining region Y)-box 10
 solute carrier family 25 (mitochondrial carrier, adenine nucleotide translocator), member 4
 transformer-2 alpha (htra-2 alpha)
 hypothetical protein FLJ22573
 KIAA0244 protein
 cholinergic receptor, nicotinic, alpha polypeptide 4
 TAK1-binding protein 2
 GABA(A) receptor-associated protein-like 2
 SSA protein SS-56
 leucine aminopeptidase
 high-mobility group (nonhistone chromosomal) protein 14
 ras homolog gene family, member C
 nuclear and coiled-body phosphoprotein 1
 UDP-Gal:betaGlcNAc 6-epimerase 1,4-galactose 4-epimerase, polypeptide 5

potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3
 SNRNP assembly defective 1 homolog
 DKFZP566C134 protein
 KIAA0141 gene product
 hypothetical protein, expressed in osteoblast
 putative mitochondrial space protein 32.1
 programmed cell death 4 (neoplastic transformation inhibitor)
 insulin-degrading enzyme
 death-associated protein 8
 KIAA0930 protein
 methyltetrahydrofolate dehydrogenase (NADP+ dependent), methyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthetase
 serologically defined breast cancer antigen 84
 CGI-43 protein
 KIAA0153 protein
 hypothetical protein FLJ120411
 dolichyl-P-Glc:Men8GlcNAc2-PP-dolichylglucosyltransferase
 asparnyglucosaminidase
 purinergic receptor P2Y, G-protein coupled, 11
 hemochromatosis
 translocated promoter region (to activated MET oncogene)
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1
 testis-specific transcript, Y-linked 1
 phosphoribosylglycinamide formyltransferase, phosphoribosylglycinamide synthetase, phosphoribosylaminoimidazole synthetase
 growth factor independent 1B (potential regulator of CDKN1A, translocated in CML)
 islet cell autoantigen 1 (69kD)
 gamma-aminobutyric acid (GABA) A receptor, alpha 5
 vesicle trafficking, beach and anchor containing
 transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase)
 CGI-02 protein
 Consensus includes gb:BE8986549 /FEA+EST /DB_XREF=gi:10369881 /DB_XREF=est:7164b02.x1 /CLONE=IMAGE:3299403 /UG=Hs 44131 KIAA0974 protein
 apoptosis inhibitor
 melanoma antigen, family B, 4
 RAB3 GTPase-ACTIVATING PROTEIN
 tropomyosin 1 (alpha)
 upstream transcription factor 2, c-fos interacting
 GPAA1P anchor attachment protein 1 homolog (yeast)
 hypothetical protein similar to mouse aldolase reductase 6 (renal)
 tetraarcopeptide repeat domain 3
 fem-1 homolog b (C. elegans)
 DKFZP568L0724 protein
 KIAA1097 protein
 hypothetical protein FLJ10658
 gamma-glutamyltransferase-like 3
 Consensus includes gb:AK026877.1 /DEF=Homo sapiens cDNA: FLJ23324 fts, clone HEP12482, highly similar to HUMMYOHC8 Human nonmuscle myosin heavy chain-B (MN
 KIAA0418 gene product
 tau-like kinase 2
 U6 snRNA-associated Sm-like protein LSM8
 ceroid-lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)
 hypothetical protein
 KIAA0901 protein
 deleted in lung and esophageal cancer 1
 hypothetical protein FLJ10422
 heterogeneous nuclear ribonucleoprotein A/B
 up-regulated by BCG-CWS
 hexosaminidase A (alpha polypeptide)
 hypothetical protein
 dihydrofolate reductase
 ZAP70 protein
 leukocyte receptor cluster (LRC) member 4
 potassium voltage-gated channel, shaker-related subfamily, beta member 1
 95 kDa neuroblastoma protein binding protein
 E2F transcription factor 1
 eukaryotic translation initiation factor 3, subunit 10 (theta, 150/170kD)

NM_002249 1.36
 NM_006590 1.36
 AF161402 1.36
 NM_014773 1.36
 NM_006820 1.36
 AF050198 1.36
 NM_014456 1.36
 NM_004969 1.36
 NM_001350 1.36
 AK025608 1.36
 NM_005956 1.36
 AF091085 1.36
 NM_016011 1.36
 BF965437 1.36
 AB050442 1.36
 NM_013339 1.36
 NM_000027 1.36
 NM_002566 1.36
 AF115264 1.36
 NM_003292 1.36
 AL040633 1.36
 AF000990 1.36
 AF008655 1.36
 NM_004188 1.36
 NM_004968 1.36
 AF061785 1.36
 W60686 1.36
 BC003551 1.36
 NM_012123 1.36
 BE856549 1.36
 NM_021831 1.36
 NM_002367 1.36
 AK022494 1.36
 NM_000366 1.36
 NM_003367 1.36
 NM_003801 1.36
 AL096767 1.36
 AU131711 1.36
 AW139179 1.36
 AU158148 1.36
 AB029020 1.36
 NM_018248 1.36
 AL049709 1.36
 AL042697 1.36
 NM_014631 1.36
 BF112255 1.35
 NM_016200 1.35
 AA602532 1.35
 NM_019083 1.35
 AB011173 1.35
 AB60341 1.35
 NM_018091 1.35
 NM_004499 1.35
 AL049963 1.35
 AL523158 1.35
 AF151022 1.35
 BC000192 1.35
 AB92986 1.35
 BC006309 1.35
 U16953 1.35
 AI379492 1.35
 NM_005225 1.35
 BE614908 1.35

hypothetical protein FLJ10774	NM_024662	1.35
Consensus includes gb:W84525 /FEA=EST /DB_XREF=gi:1355637 /DB_XREF=est:zd90q05.s1 /CLONE=IMAGE:356800 /UG=Hs.7949 DKFZP586B2420 protein	W84525	1.35
G protein-binding protein CRFG	NM_012341	1.35
heat shock 70kD protein 4	AA043348	1.35
Sney homolog, Y chromosome (mouse)	NM_004653	1.35
tumor protein, translationally-controlled 1	AI886178	1.35
protein kinase C, α	L18964	1.35
DEAD/HD (Asp-Glu-Ala-Asp/His) box polypeptide 9 (RNA helicase A, nuclear DNA helicase II; leukophycin)	BE561014	1.35
phosphoglucomutase 1	NM_002633	1.35
CDI-203 protein	NM_020408	1.35
Consensus includes gb:AI348006 /FEA=EST /DB_XREF=gi:4085212 /DB_XREF=est:qp61b05.x1 /CLONE=IMAGE:1927473 /UG=Hs.469 succinate dehydrogenase complex, β	AI348006	1.35
NICE-5 protein	NM_017582	1.35
sentrin/SUMO-specific protease 3	NM_015670	1.35
cyclin-dependent kinase 9 (CDC2-related kinase)	NM_001261	1.35
protein phosphatase 1, regulatory (inhibitor) subunit 2	AW614578	1.35
hypothetical protein DT1P1A10	AA223871	1.35
Bardet-Biedl syndrome 4	AI813772	1.35
protein tyrosine phosphatase type IVA, member 2	U48297	1.35
hypothetical protein	U69190	1.35
Spt-B transcription factor (Spi-1/PU.1 related)	NM_003121	1.35
zinc finger protein 43 (a Kruppel-associated box (KRAB) domain polypeptide)	NM_003425	1.35
heat shock factor binding protein 1	AK026575	1.35
hypothetical protein FLJ22622	NM_025151	1.35
Cluster Inc. AI952879:wd24c06.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2508394 /clone_end=3 /gb=AI962879 /gi=5755592 /ug=Hs.81920 /len=490	AI962879	1.35
H2B histone family, member L	NM_003526	1.35
Consensus includes gb:BG031677 /FEA=EST /DB_XREF=gi:12422203 /DB_XREF=est:602300830F1 /CLONE=IMAGE:4402326 /UG=Hs.58210 ESTs	BG031677	1.35
hypothetical protein FLJ22073	NM_024827	1.35
soluble carrier family 25 (mitochondrial carrier; adenine nucleotide translocator), member 5	NM_017654	1.35
potassium channel modulator/factor	NM_001152	1.35
SAR1 protein	NM_020122	1.35
serine/threonine kinase 24 (STE20 homolog, yeast)	NM_003940	1.35
MCM4 mitochromosome maintenance deficient 4 (S. cerevisiae)	NM_020150	1.35
hypothetical protein FLJ11937	AF083420	1.35
OneJ (Hsp40) homolog, subfamily A, member 3	AI936566	1.35
pro-platelet basic protein (includes platelet basic protein, beta-thromboglobulin, connective tissue-activating peptide III, neutrophil-activating peptide-2)	BF575213	1.35
Consensus includes gb:AI798823 /FEA=EST /DB_XREF=gi:5364295 /DB_XREF=est:wb3b05.x1 /CLONE=IMAGE:2348625 /UG=Hs.1050 plectstrin homology, Sec7 and coiled coil	NM_022765	1.35
ves 2 oncogene	NM_005147	1.35
hypothetical protein MGC4022	R64130	1.35
amyloid beta (A4) precursor protein-binding, family B, member 1 (Fe65)	NM_003371	1.35
lactate dehydrogenase B	BF566155	1.35
ecoonucleoside triphosphate diphosphohydrolase 6 (putative function)	NM_001164	1.35
ribosomal protein S4, X-linked	BE042354	1.35
methy-CpG binding protein 2 (Rett syndrome)	L37288	1.35
ubiquitin specific protease 14 (IRNA-guanine transglycosylase)	NM_001247	1.35
protein phosphatase 3 (formerly 2B), catalytic subunit, beta isoform (calcineurin A beta)	AL035603	1.35
protein tyrosine kinase 9	AI913365	1.35
nuclear protein ANK1	NM_005151	1.35
peroxisomal acyl-CoA thioesterase 1	NM_021132	1.35
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	AW665024	1.35
1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)	NM_018454	1.35
potassium voltage-gated channel, shaker-related subfamily, member 6	NM_005469	1.35
hypothetical protein	AV725365	1.35
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 42kD	U56417	1.35
tyrosyl-DNA phosphodiesterase	AW134812	1.34
regulator of Fas-induced apoptosis	AI926407	1.34
ATPase, Na ⁺ /K ⁺ transporting, alpha 1 polypeptide	BC005863	1.34
hypothetical protein from EUROIMAGE 1987170	BF034973	1.34
glutathione transferase zeta 1 (maleylacetoacetate isomerase)	NM_018319	1.34
	AI084226	1.34
	NM_000701	1.34
	AL136925	1.34
	BC001453	1.34

SET domain, bifurcated 1
 Consensus includes gb:AL024509 /DEF=Human DNA sequence from clone 522P13 on chromosome 6p21.31-22.3. Contains a 60S Ribosomal Protein L21 pseudogene and a hypoxia-inducible factor 1, alpha subunit inhibitor
 metallothionein 1G
 aldo-keto reductase family 1, member C1 (dihydrodiol dehydrogenase 1; 20-alpha (beta)-hydroxysteroid dehydrogenase)
 immunoglobulin kappa variable 1J015-118
 Consensus includes gb:A071147 /FEA=EST /DB_XREF=gi:4851478 /DB_XREF=IMAGE:2297218 /UG=Hs.81360 Homo sapiens cDNA: FLJ121927 fr. A071147
 plectrin
 ribosomal protein S7
 pituitary tumor-transforming 1
 Consensus includes gb:AA650558 /FEA=EST /DB_XREF=gi:2577886 /DB_XREF=IMAGE:1192002 /UG=Hs.325202 ESTs, highly similar to GBAS_H1, AA650558
 solute carrier family 2 (facilitated glucose transporter), member 1
 hypothetical protein FLJ21839
 cleavage stimulation factor, 3' pre-RNA, subunit 3, 77kD
 tetranucleotide repeat domain 4
 RAB2, member RAS oncogene family
 erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-linked)
 inhibitor of DNA binding 4, dominant negative helix-loop-helix protein
 hypothetical protein FLJ20827
 fibroblast growth factor 5
 Consensus includes gb:AX022663.1 /DEF=Homo sapiens cDNA FLJ12601 f1s, clone NT2RM4001414, /FEA=mRNA /DB_XREF=gi:10434187 /UG=Hs.288549 Homo sapiens c. AK022663
 lamin B1
 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5 (16kD, SGDh)
 chromodomain protein, Y chromosome-like
 iniquitous homeobox protein 5
 nuclear factor (erythroid-derived 2)-like 1
 aryl hydrocarbon receptor nuclear translocator
 Consensus includes gb:AL133173 /DEF=Human DNA sequence from clone RP11-453N3 on chromosome 10 Contains the 3' end of a gene similar to ABCD1 (ATP-binding cassette AL133173)
 hypothetical protein MGC11308
 carbonic anhydrase II
 mannosidase, alpha, class 2B, member 1
 ribosomal protein L34
 Consensus includes gb:AL109622 /DEF=Human DNA sequence from clone RP3-526F5 on chromosome Xq26.3-28. Contains the gene for a novel protein similar to UBE2N (ut S55735)
 immunoglobulin heavy constant mu
 ras-like protein
 growth factor receptor-bound protein 2
 KIAA1449 protein
 ubiquitin-conjugating enzyme E2 variant 1
 FLN29 gene product
 Rho guanine exchange factor (GEF) 12
 Cluster Ind. N32782:yy91903.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:259638 /clone_end=3 /gb=N32782 /gi=1153181 /ug=Hs.13138 /len=668
 solute carrier family 35 (CMP-sialic acid transporter), member 1
 peroxisomal biogenesis factor 3
 sine oculis homeobox homolog 2 (Drosophila)
 hypothetical protein FLJ12525
 second mitochondria-derived activator of caspase
 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit b, isoform 1
 putative zinc finger protein from EUROMIMAGE 566589
 aspartyl aminopeptidase
 hypothetical protein FLJ12442
 collagen
 Consensus includes gb:AL049987.1 /DEF=Homo sapiens mRNA: cDNA DKFZ564F112 (from clone DKFZ564F112), /FEA=mRNA /DB_XREF=gi:4884238 /UG=Hs.168361 H.AL049987
 hypothetical protein FLJ22477
 glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)
 hypothetical protein FLJ12847
 H3PC166 protein
 H3PC166 protein
 similar to Caenorhabditis elegans protein C42C1.9
 gogli associated, gamma adaptin ear containing, ARF binding protein 2
 protocadherin beta 17 pseudogene
 hypothetical protein FLJ20550
 general transcription factor IIF, polypeptide 2 (44kD subunit)
 breast cancer anti-estrogen resistance 3

Consensus includes gb:AL359954 /DEF=Human DNA sequence from clone RP11-318P23 on chromosome 20 Contains a TAR DNA-binding protein pseudogene, ESTs, STSs. AL359954
Consensus includes gb:AF070537.1 /DEF=Homo sapiens clone 24606 mRNA sequence. /FEA=mRNA/DB_XREF=gi:3387894 /UG=Hs.17481 Homo sapiens clone 24606 mRt AF070537
equilene epoxide 1.33
AF098865 1.33
CDP-diacylglycerol synthase (phosphatidate cytidylyltransferase) 2 1.33
AF56982 1.33
hepatocyte growth factor-regulated tyrosine kinase substrate 1.33
AF260568 1.33
AFG3 ATPase family gene 3-like 2 (yeast) 1.33
X71810 1.33
retinoblastoma binding protein 4 1.33
A5566082 1.33
syntaxin 7 1.33
M74220 1.33
clastinogen 1.33
NM_018099 1.33
hypothetical protein FLJ10462 1.33
growth differentiation factor 1 1.33
NM_001492 1.33
Cluster Ind. AA601208:nc14c12.1 Homo sapiens cDNA. 3' end /clone=IMAGE:1100662 /clone_end=3 /gb=AA601208 /gi=2434833 /ug=Hs.220255 /len=488
early development regulator 2 (homolog of polyhomeotic 2) 1.33
AA601208 1.33
hypothetical protein FLJ12528 1.33
NM_004427 1.33
gb:NM_005703.2 /DEF=Homo sapiens upstream regulatory element binding protein 1 (UREB1), mRNA. /FEA=CDS /GEN=UREB1 /PROD=upstream regulatory element binding protein 1 (UREB1), mRNA. /FEA=mRNA /DB_XREF=gi:4884400 /UG=Hs.225981 AL050185
broadly acidic repeat containing 3 1.33
NM_005703 1.33
broadly acidic repeat containing 3 1.33
AL541302 1.33
BUP protein 1.33
NM_012071 1.33
hypothetical protein FLJ20360 1.33
NM_017782 1.33
hypothetical protein FLJ22594 1.33
NM_024525 1.33
creatine kinase, muscle 1.33
NM_001824 1.33
BC002711 1.33
cell division cycle 42 (GTP binding protein, 25kD) 1.33
AL527773 1.33
active BCR-related gene 1.33
A630178 1.33
hypothetical protein DKFZp762A227 1.33
transposon-derived Buser3 transposase-like 1.33
HSPC022 protein 1.33
NM_022090 1.33
hypothetical protein FLJ11506 1.33
NM_024666 1.33
ubiquitin specific protease 16 1.33
NM_006447 1.33
membrane-associated tyrosine- and threonine-specific cdc2-inhibitory kinase 1.33
NM_004203 1.33
hypothetical protein FLJ11848 1.33
NM_025155 1.33
interferon, alpha 1 1.33
NM_024013 1.33
thymosin, beta 4, Y chromosome 1.33
NM_007062 1.33
nuclear phosphoprotein similar to S. cerevisiae PWP1 1.33
AW151887 1.33
heparan sulfate 2-O-sulfotransferase 1.33
AA918442 1.33
insulin-degrading enzyme 1.33
NM_000999 1.33
ribosomal protein L38 1.33
AA961748 1.33
ribosomal protein L13 1.33
NM_022744 1.33
hypothetical protein FLJ13868 1.33
DKFZP564O092 protein 1.33
A1269117 1.33
hypothetical protein 1.33
NM_015702 1.33
interleukin 12 receptor, beta 1 1.33
NM_005535 1.33
vimentin 1.33
A922599 1.33
ubiquitin-conjugating enzyme E2 variant 1 1.33
BG164064 1.33
FliJ homolog 1 (E. coli) 1.33
AV723177 1.33
zincfin 1.33
NM_013403 1.33
dynactin 2 (p50) 1.33
NM_006400 1.33
phospholipase C, beta 2 1.33
NM_004573 1.33
ION3 protein 1.33
AW339587 1.33
NY-REN-24 antigen 1.33
cytochrome b-561 1.33
A1200896 1.33
ras homolog gene family, member 1.33
AW003733 1.33
phosphatidylinositol glycan, class N 1.33
E2F transcription factor 2 1.33
NM_012327 1.33
chromosome 5 open reading frame 8 1.33
NM_004091 1.33
Consensus includes gb:AL050185.1 /DEF=Homo sapiens mRNA, cDNA DKFZp586A0423 (from clone DKFZp586A0423). /FEA=mRNA /DB_XREF=gi:4884400 /UG=Hs.225981 AL050185
putative DNA binding protein 1.33
AL523380 1.33
ribosomal protein S2 1.33
BF125158 1.33
centaurin, beta 1 1.33
NM_014716 1.33
myosin, light polypeptide, regulatory, non-sarcomeric (20kD) 1.33
NM_006471 1.33
KIAA0385 gene product 1.33
amyloid beta (A4) precursor protein-binding, family B, member 2 (Fe65-like) 1.33
BG252853 1.33
TNFRSF1A-associated via death domain 1.33
U82325 1.33
L41690 1.33
interferon gamma receptor 1 1.33
NM_000416 1.33
centaurin, alpha 1 1.33
AW050627 1.33

poliovirus receptor-related 1 (herpesvirus entry mediator C; nectin)
 H3 histone, family 3B (H3.3B)
 KIAA0998 protein
 chromosome 11 open reading frame 23
 zinc finger protein, 103 homolog (mouse)
 hypothetical protein FLJ21865
 RNA helicase family
 nuclear antigen Sp100
 fragile X mental retardation 2
 AHNK nucleoprotein (desmoyotin)
 hypothetical protein FLJ20045
 Consensus includes gb:AU154486 /FEA=EST /DB_XREF=gi:11016007 /DB_XREF=est:AU154486 /CLONE=NT2RP4001238 /UG=Hs.119023 SMC2 (structural maintenance of Nt protein 2)
 H3 histone family, member 1
 hypothetical protein dJ465N24.2.1
 hypothetical protein FLJ22246
 RNA guanylyltransferase and 5'-phosphatase
 integral membrane protein 3
 copine 1
 thyroid hormone receptor interactor 4
 vesicular protein sorting 45B (yeast)
 IGF-II mRNA-binding protein 3
 ubiquitin-conjugating enzyme E2L 3
 hypothetical protein DKFZ6564D0478
 hypothetical protein FLJ10583
 enocephalin
 ribosomal protein L31
 TERF1 (TRF1)-interacting nuclear factor 2
 Cipl-1-interacting zinc finger protein
 proteasome (prosome, macropain) 26S subunit, ATPase, 2
 oysteroid binding protein-like 3
 Consensus includes gb:BF445142 /FEA=EST /DB_XREF=gi:11510203 /DB_XREF=est:rad21b05.x1 /CLONE=MAGE:3366081 /UG=Hs.797 nuclear transcription factor Y, alpha
 Consensus includes gb:AK025343.1 /DEF=Homo sapiens cDNA: FLJ21690 fs, clone COL09538, /FEA=mRNA /DB_XREF=gi:10437841 /UG=Hs.306802 Homo sapiens cDNA: AK022874
 DKFZP596E1519 protein
 tumor protein, translationally-controlled 1
 calpain 10
 CLIP-associated protein 2
 BRCA1 associated protein-1 (ubiquitin carboxy-terminal hydrolase)
 tubulin-specific chaperone 8
 chromosome 11 hypothetical protein ORF4
 splicing factor 3b, subunit 1, 155kD
 hypothetical protein FLJ20721
 chorionic somatomammotropin hormone 1 (placental lactogen)
 paraneoplastic antigen MA1
 protein-rich protein with nuclear targeting signal
 hypothetical protein DKFZp34E1723
 STIP1 homology and U-Box containing protein 1
 calcium/calmodulin-dependent protein kinase (CaM kinase) II beta
 BCL2-associated X protein
 low density lipoprotein receptor-related protein 8, apolipoprotein e receptor
 KIAA0690 protein
 zinc finger protein 277
 hypothetical protein FLJ13910
 UBX domain-containing 2
 protease, serine, 22
 ubiquitin specific protease 4 (proto-oncogene)
 KIAA0978 protein
 putative protein kinase NY-REN-64 antigen
 hypothetical protein DKFZp434K0920
 adaptor-related protein complex 4, sigma 1 subunit
 muS homolog 6 (E. coli)
 KIAA0372 gene product
 tripartite motif-containing 36

NM_002855 1.32
 AW138159 1.32
 AB023215 1.32
 NM_018312 1.32
 NM_005667 1.32
 AL044087 1.32
 AA156961 1.32
 NM_003113 1.32
 NM_002025 1.32
 BG287862 1.32
 BC002894 1.32
 NM_020202 1.32
 NM_021016 1.32
 AF247168 1.32
 NM_025232 1.32
 AB009023 1.32
 NM_030926 1.32
 NM_003915 1.32
 NM_016213 1.32
 AF165513 1.32
 AL023775 1.32
 AI819709 1.32
 AL136883 1.32
 NM_018148 1.32
 NM_001988 1.32
 NM_000993 1.32
 NM_012461 1.32
 NM_012127 1.32
 NM_002803 1.32
 AI202969 1.32
 BF445142 1.32
 AK025343 1.32
 AK022874 1.32
 AL565449 1.32
 NM_023089 1.32
 AI741784 1.32
 NM_004656 1.32
 NM_004607 1.32
 AA813184 1.32
 AI739389 1.32
 NM_017941 1.32
 NM_022641 1.32
 NM_006029 1.32
 AF279899 1.32
 AI743331 1.32
 NM_005861 1.32
 AF112472 1.32
 U19599 1.32
 NM_004631 1.32
 AK021460 1.32
 NM_021994 1.32
 BF671894 1.32
 AU149808 1.32
 AI446043 1.32
 BE256900 1.32
 NM_016123 1.32
 NM_017590 1.32
 AF155159 1.32
 NM_000179 1.32
 BE566023 1.32
 NM_018700 1.32

3-phosphoinositide dependent protein kinase-1
 DKFZP568I2223 protein
 primase, polypeptide 2A (58kd)
 Consensus includes gb:AY007126.1 /IDEF=Homo sapiens clone CDABP0028 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3956024 /UG=Hs.24435 Homo sapiens clone CDA AY007126
 KIAA0692 protein
 myotubularin related protein 2
 nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, alpha
 Consensus includes gb:BF057059 /FEA=EST /DB_XREF=gi:10810955 /DB_XREF=est:7k15c07.x1 /CLONE=IMAGE:3444468 /UG=Hs.12421 hypothetical protein
 microtubule-associated proteins 1A/1B light chain 3
 sense domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6C
 9-adenosylhomocysteine hydrolase-like 1
 v-ets erythroblastosis virus E26 oncogene homolog 2 (avian)
 kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)
 bone marrow stromal cell antigen 1
 Consensus includes gb:BG429214 /FEA=EST /DB_XREF=gi:13335720 /DB_XREF=est:802498083f1 /CLONE=IMAGE:4611836 /UG=Hs.258144 zinc finger protein 33b (KIX)
 biphenyl hydrolase-like (serine hydrolase; breast epithelial mucin-associated antigen)
 hypothetical protein
 zinc finger protein 143 (clone pHz-1)
 upstream transcription factor 2, c-fos interacting
 hypothetical protein MGC3067
 hypothetical protein MGC2478
 hypothetical protein FLJ20452
 sense domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4D
 Consensus includes gb:AK025177.1 /IDEF=Homo sapiens cDNA: FLJ21524 fts, clone COL05921. /FEA=mRNA /DB_XREF=gi:10437640 /UG=Hs.306778 Homo sapiens cDNA: AK025177
 hypothetical protein FLJ12595
 Consensus includes gb:BE538424 /FEA=EST /DB_XREF=gi:3767069 /DB_XREF=est:801088256f1 /CLONE=IMAGE:3454693 /UG=Hs.288283 Homo sapiens cDNA: FLJ2231 BE538424
 translocase of inner mitochondrial membrane 44 homolog (yeast)
 pumilio homolog 1 (Drosophila)
 nuclear receptor coactivator 4
 mult. homolog 1, colon cancer, nonpolyposis type 2 (E. coli)
 Sec23-interacting protein p125
 G protein-coupled receptor 27
 hypothetical protein FLJ20608
 kionate 2-sulfatase (Hunter syndrome)
 tubulin, beta polypeptide
 tumor rejection antigen (gp96) 1
 SKI-interacting protein
 G protein-coupled receptor 43
 endocrine regulator
 VRK3 for vaccinia related kinase 3
 protein phosphatase 4, regulatory subunit 1
 Consensus includes gb:AW128846 /FEA=EST /DB_XREF=gi:5116784 /DB_XREF=est:xd46106.x1 /CLONE=IMAGE:2596635 /UG=Hs.244737 ESTs. Weakly similar to neurons
 hypothetical protein FLJ11209
 Consensus includes gb:BC004409.1 /IDEF=Homo sapiens, clone IMAGE:3638994. mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3638994) /DB_XREF=BC004409
 small nuclear ribonucleoprotein D3 polypeptide (18kd)
 KIAA0848 protein
 RAB4B, member RAS oncogene family
 RAP1, GTP-GDP dissociation stimulator 1
 hypothetical protein MGC2875
 KIAA0810 protein
 protein tyrosine phosphatase type IVA, member 2
 salivase 3 (membrane salivase)
 ubiquitin C-terminal hydrolase UCH37
 glutamate-cysteine ligase, catalytic subunit
 C2H2 (Krueppel-type) zinc finger protein
 stannin
 prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)
 PAS-serine/threonine kinase
 5-hydroxytryptamine (serotonin) receptor 2C
 regulator of G-protein signalling 14
 paired-like homeodomain transcription factor 1
 PAL-1 mRNA-binding protein
 ribosomal protein L24

SRY (sex determining region Y)-box 4	A189477	1.31
perlecanin 2 (kndrin)	NM_006031	1.31
SBB126 protein	NM_018846	1.31
calpastatin	AF327443	1.31
cleavage and polyadenylation specific factor 6, 68kD subunit	AU149387	1.31
paritin	D88862	1.31
hypothetical protein FLJ12686	NM_024595	1.31
2N10 homolog, centrosomal/retinoblastoma protein (Drosophila)	NM_004724	1.31
hypothetical protein FLJ10057	NM_017684	1.31
hypothetical protein FLJ21657	NM_022463	1.31
G protein-coupled receptor 30	U63917	1.31
vesicular protein sorting 35 (yeast)	NM_018206	1.31
solute carrier family 18 (vesicular monoamine), member 2	AZ69290	1.31
acrosomal vesicle protein 1	NM_020110	1.31
ATP synthase, H+ transporting, mitochondrial F1 complex, O subunit (oligomycin sensitivity conferring protein)	S77356	1.31
cytochrome c oxidase subunit Va	NM_004255	1.31
PPTAIRE protein kinase 1	NM_012395	1.31
EGF-like domain, multiple 3	AL134303	1.31
hypothetical protein FLJ10280	NM_018047	1.31
heat shock transcription factor 4	AB029348	1.31
adenylyl cyclase-associated protein	AA806142	1.31
KIAA0876 protein	NM_015043	1.31
KIAA0873 protein	AB014573	1.31
mudS homolog 3 (E. coli)	NM_002439	1.31
DKFZP564J102 protein	AL080065	1.31
KIAA1232 protein	AB033056	1.31
zinc finger protein 212	NM_012256	1.31
Consensus includes gb:A1721229 /FEA=EST /DB_XREF=gi:5038485 /DB_XREF=est:as68c10.x1 /CLONE=IMAGE:2333874 /UG=Hs.326456 hypothetical protein FLJ20030	A1721229	1.31
tumoral alkaline phosphatase	NM_000143	1.31
tenascin 1	NM_004109	1.31
transcriptional co-activator with PDZ-binding motif (TAZ)	BF674349	1.31
squamous cell carcinoma antigen recognised by T cells 3	AW173076	1.31
hypothetical protein MDS018	NM_021823	1.31
exosome component Rps41	A1571288	1.31
glucocorticoid receptor DNA binding factor 1	AB070100	1.31
gb:AB056895.1 /DEF=Homo sapiens GHRHRp sv mRNA for growth hormone releasing hormone receptor pituitary splice variant, complete cds. /FEA=mRNA /GEN=GHRHRp	NM_005885	1.31
lactotransferrin	NM_005762	1.31
tripartite motif-containing 28	W07773	1.31
Cluster Ind. W07773:zb03g04.1 Homo sapiens cDNA, 5' end /clone=IMAGE:301014 /clone_end=5' gb=W07773 /gi=1281785 /ug=Hs.9739 /len=533	W07773	1.31
HEMK homolog 7kb	AF131220	1.31
cholesterol oxidase	AB023203	1.31
hypothetical protein MGC3121	AA161130	1.31
olfactory receptor, family 2, subfamily 5, member 2	NM_019897	1.31
uncharacterized hypothalamus protein HARP11	AK022248	1.30
KIAA0887 protein	NM_014938	1.30
CGI-148 protein	NM_016076	1.30
gene with multiple splice variants near HD locus on 4p16.3	NM_003704	1.30
MAOS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	AL536517	1.30
Consensus includes gb:AL390857 /DEF=Human DNA sequence from clone RP11-51N22 on chromosome 13 Contains ESTs, STSs and GSSs. Contains an HNRPA1 (hetero	AL390857	1.30
secreted fuzed-related protein 1	A132407	1.30
hypothetical protein FLJ10315	NM_018056	1.30
phosphatidylinositol binding dalphin assembly protein	AV721177	1.30
CD19 antigen	NM_001770	1.30
vesicle-associated membrane protein 4	NM_003762	1.30
MZ7530 Human 26S ribosomal RNA gene, complete cds (5', 3' represent transcript regions 5 prime, Middle, and 3 prime respectively)	M27830	1.30
suppressor of var1, 3-like 1 (S. cerevisiae)	NM_003171	1.30
serine/threonine kinase 11 (Peutz-Jeghers syndrome)	AF035625	1.30
synovial sarcoma, X breakpoint 2	AU152088	1.30
KIAA0332 protein	NM_003147	1.30
procadherin beta 13	NM_018933	1.30
colony stimulating factor 2 receptor, beta, low-affinity (granulocyte-macrophage)	AV756141	1.30
cAMP responsive element binding protein 1	M34356	1.30
chromosome 2 open reading frame 2	NM_019053	1.30

Consensus includes gb:J04742 /DEF=Human autonomous replicating sequence H1 (ARSH1) /FEA=CDS /DB_XREF=gi:179028 /UG=Hs.247345 Human autonomous replicatin J04742
KJAA0090 protein
Consensus includes gb:AK002078.1 /DEF=Homo sapiens cDNA FLJ11214 f1s, clone PLACE1007990. /FEA=mRNA /DB_XREF=gi:7023738 /UG=Hs.281618 Homo sapiens cD AK002078
DNFZP564C196 protein
inositol polyphosphate phosphatase-like 1
glycosylphosphatidylinositol specific phospholipase D1
hypothetical protein FLJ22855
BTB (POZ) domain containing 1
hypothetical protein FLJ20343
Consensus includes gb:AK22005 /DEF=Human interleukin 2 gene, clone pATlacIL-2C2TT, complete cds, clone pATlacIL-2C2TT /FEA=CDS /DB_XREF=gi:186300 /UG=Hs.2471422005
zinc finger protein 36, C3H type-like 1
nuclear transport factor 2 (placental protein 15)
hypothetical protein MGC2483
zinc finger protein 278
fodilin 1
HSPC035 protein
Consensus includes gb:AL109942 /DEF=Human DNA sequence from clone RP3-47J18 on chromosome 6p25.3-26 Contains 3 end of the MAP3K4 (mitogen-activated protein
Cluster loc. A424872 /g3810.11 Homo sapiens cDNA, 3 end /clone=IMAGE-2111083 /clone_end=3 /gb=A424872 /gi=4270895 /ug=Hs.159375 /len=443
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 10 (RNA helicase)
gpgi associated, gamma adspilin ear containing, ARP binding protein 2
hypothetical protein MGC355
Cluster loc. AW028481 /w14a04.x1 Homo sapiens cDNA, 3 end /clone=IMAGE-990913 /clone_end=3 /gb=AW028481 /gi=5880011 /ug=Hs.48396 /len=496
alanine-glyoxylate aminotransferase (oxalosis 1; hyperoxaluria 1; glycolicaciduria; serine-pyruvate aminotransferase)
origin recognition complex, subunit 1-like (yeast)
Ras9 effector p40
phosphofructokinase, liver
zinc finger protein 16 (K0X 9)
nucleix (nucleoside diphosphate linked moiety X)-type motif 2
metallothionein IV
antigen identified by monoclonal antibody Ki-67
arsenate resistance protein ARS2
RAS guanyl releasing protein 2 (calcium and DAG-regulated)
glucose-6-phosphate dehydrogenase
KIAA0370 protein
hypothetical protein FLJ20783
nuclear transcription factor Y, alpha
prostate epithelium-specific Ets transcription factor
HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae)
ribosomal protein L7
hypothetical protein FLJ21840
Rho GTPase activating protein 1
hypothetical protein similar to preferentially expressed antigen of melanoma
solute carrier family 9 (sodium/hydrogen exchanger), isoform 3 regulatory factor 1
CGI-30 protein
glycerol-3-phosphate dehydrogenase 2 (mitochondrial)
lysine 3-monooxygenase/tyrosine 5-monooxygenase activation protein, epsilon polypeptide
Consensus includes gb:L24521.1 /DEF=Human (transformation-related protein mRNA, 3 end, /FEA=mRNA /PROD=transformation-related protein /DB_XREF=gi:403459 /UG=H L24521
nombold, veiriel-like 1 (Drosophila)
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4
KIAA0046 protein
nuclear receptor co-repressor 2
chloride channel, calcium activated, family member 4
mannosidase, alpha, class 1B, member 1
heparin-binding growth factor binding protein
KIAA0063 gene product
fumarate hydratase
sorting nexin 11
serine/threonine kinase 18
DHHC1 protein
v-myo myeloblastosis viral oncogene homolog (avian)-like 1
protein tyrosine kinase 2 beta
small nuclear ribonucleoprotein 70kD polypeptide (RNP antigen)
acidic protein rich in leucines

folate receptor 3 (gamma)	NM_000804	1.29
bedin 1 (coiled-coil, myosin-like BCL2 interacting protein)	NM_003766	1.29
hypothetical protein MGC4825	AF061264	1.29
dyxerlin, limo girle muscular dystrophy 2B (autosomal recessive)	NM_003494	1.29
patently expressed 10	AL582636	1.29
surfeit 1	NM_003172	1.29
regulator of G-protein signalling 16	BF304996	1.29
regucalcin (senescence marker protein-30)	D31815	1.29
fatty-acid-Coenzyme A ligase, long-chain 2	NM_021122	1.29
granulin	BC000324	1.29
similar to yeast Upt3, variant B	NM_023010	1.29
gamma-glutamyl carboxylase	BE326952	1.29
OnaJ (Hsp40) homolog, subfamily C, member 4	NM_005528	1.29
Interleukin-1 receptor-associated kinase 1	NM_001569	1.29
hypothetical protein FLJ10719	BG403815	1.29
aldo-keto reductase family 1, member C1 (dihydrodiol dehydrogenase)	S68280	1.29
rTS beta protein	NM_017512	1.29
transducin (beta)-like 2	AI608789	1.29
pre-B-cell colony-enhancing factor	NM_005746	1.29
solute carrier family 2B (nucleoside transporters), member 1	AF078117	1.29
karyopherin (importin) beta 1	AA861608	1.29
selenoprotein X, 1	NM_016332	1.29
hypothetical protein FLJ10688	AF062733	1.29
natural killer-tumor recognition sequence	AI688640	1.29
EH-domain containing 1	AF001434	1.29
filamin B, beta (actin binding protein 278)	M62994	1.29
RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae)	NM_002875	1.29
LamC (bacterial lamibiotic synthetase component C)-like 2	NM_018697	1.29
v-naf musculoskeletal fibrosarcoma oncogene homolog F (avian)	NM_012323	1.29
protein kinase C, alpha	NM_002737	1.29
protein necrosis factor, alpha-induced protein 3	NM_006280	1.29
mannosyl (alpha-1,6)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase	NM_006280	1.29
Consensus includes gb:A149535 /FEA=EST /DB_XREF=gi:3678004 /DB_XREF=est:3678004 /CLONE=IMAGE:1714989 /UG=Hs.24064 Homo sapiens mRNA; cDNA DKFZ	AW103285	1.29
KIAA0802 protein	AB018345	1.29
phosphotransferase, liver	NM_002826	1.29
gb:BC001685.1 /DEF=Homo sapiens, similar to cadherin 1, type 1, E-cadherin (epithelial), done MGC:1151, mRNA, complete cds. /FEA=mRNA /PROD=Similar to cadherin 1	BC001865	1.29
hypothetical protein d434014.3	NM_025228	1.29
Consensus includes gb:BF037823 /FEA=EST /DB_XREF=gi:10746209 /DB_XREF=est:601461994F /CLONE=IMAGE:3865487 /UG=Hs.9883 OnaJ (Hsp40) homolog, subfam	BF037823	1.29
hypothetical protein FLJ13096	NM_025000	1.29
serine/arginine repetitive matrix 2	AI655799	1.29
biliverdin reductase A	AA740186	1.29
dystroglycan 1 (dystrophin-associated glycoprotein 1)	NM_004393	1.29
N-acetylneuraminic acid phosphate synthase; sialic acid synthase	NM_018946	1.29
alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, S. cerevisiae)	U09820	1.29
survival of motor neuron 1, telomeric	NM_000344	1.29
platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog)	AU150748	1.29
tripartite motif-containing 26	NM_003449	1.29
transcription factor 8 (represses interleukin 2 expression)	U12170	1.29
proteasome (prosome, microgamin) subunit, alpha type, 3	NM_002788	1.29
solute carrier family 7 (cationic amino acid transporter, y ⁺ system), member 8	AL365347	1.29
ALEX1 protein	NM_016608	1.29
leukocyte-IRNA synthetase	NM_013417	1.29
KIAA0908 protein	AA502912	1.29
mitochondrial ribosomal protein S31	AI937794	1.29
NAD, mothers against decapentaplegic homolog 2 (Drosophila)	NM_005901	1.29
cyclin M1	NM_020348	1.29
transcription elongation factor A (SII), 1	X57198	1.29
KIAA0737 gene product,	BE783632	1.29
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1	U80918	1.29
ER to nucleus signalling 1	NM_001433	1.29
CDK3 antigen (melanoma 1 antigen)	NM_001780	1.29
complement component 8, beta polypeptide	NM_000066	1.29
angiotensin-3	NM_015985	1.29

Consensus includes gb:XB4340.1 /DEF=H.sapiens mRNA (or Ig light chain, variable region (ID:CLL001VL)). /FEA=mRNA /GEN=lg VL gene /PROD=immunoglobulin light chain X84340 1.29
 Hox and leucine-rich repeat protein 5 AF157323 1.29
 rth endonuclease II-like 1 (E. coli) AC005600 1.29
 H4 histone family, member G NM_003542 1.29
 Ras homolog enriched in brain 2 BF593727 1.28
 myotubularin related protein 4 AB014547 1.28
 goyil associated, gamma adaphin ear containing, ARF binding protein 3 AV008018 1.28
 breast carcinoma amplified sequence 2 NM_005872 1.28
 SET translocation (myeloid leukemia-associated) NM_003011 1.28
 HUS1 checkpoint homolog (S. pombe) NM_004507 1.28
 transient receptor potential channel 1 U31110 1.28
 ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function) AW194947 1.28
 AL022724 1.28
 hypothetical protein MGC12335 NM_024104 1.28
 Consensus includes gb:AK025348.1 /DEF=Homo sapiens cDNA: FLJ21695 fls, clone COL09653. /FEA=mRNA /DB_XREF=gi:10437846 /UG=Hs.308803 Homo sapiens cDNA: AK025348 1.28
 diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein) BC008466 1.28
 branched chain keto acid dehydrogenase E1, beta polypeptide (maple syrup urine disease) M55575 1.28
 ribosomal protein L22 M55575 1.28
 transcription elongation factor B (SIII), polypeptide 1 (15kD, elongin C) NM_000983 1.28
 gb:Z25432.1 /DEF=H.sapiens protein-serine/threonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405742 /FL=gb:Z25432. Z25432 1.28
 hypothetical protein, clone 24751 AF070530 1.28
 surfactant 5 NM_006752 1.28
 L38951 1.28
 karyopherin (importin) beta 1 NM_012103 1.28
 BCR downstream signaling 1 NM_017518 1.28
 Consensus includes gb:Z88751 /DEF=Human DNA sequence from PAC 56089 on chromosome 1q24.1q25. Contains prolin-like pseudogene. 60S ribosomal protein L4 pseud 298751. 1.28
 hypothetical protein FLJ22187 NM_024936 1.28
 Xq28. 2000bp sequence contig. ORF AF177272 1.28
 UDP glycosyltransferase 2 family, polypeptide B28 NM_024936 1.28
 SMARF protein BE384482 1.28
 Consensus includes gb:AK025191.1 /DEF=Homo sapiens cDNA: FLJ21538 fls, clone COL08151. /FEA=mRNA /DB_XREF=gi:10437654 /UG=Hs.306783 Homo sapiens cDNA: AK025191 1.28
 JM4 protein NM_007213 1.28
 citron (Pro-interacting, serine/threonine kinase 21) A1661768 1.28
 hypothetical protein MGC15396 BC005853 1.28
 solute carrier family 5 (sodium iodide symporter), member 5 D87920 1.28
 CDC16 cell division cycle 16 homolog (S. cerevisiae) AF164598 1.28
 MHC class I polypeptide-related sequence A NM_000247 1.28
 CDC16 cell division cycle 16 homolog (S. cerevisiae) AF164598 1.28
 KIAA1198 protein A1493119 1.28
 solute carrier family 7 (cationic amino acid transporter, y+ system), member 8 AL385343 1.28
 nitrogen-activated protein kinase kinase kinase 7 NM_003188 1.28
 manic fringe homolog (Drosophila) NM_002405 1.28
 myelin transcription factor 1 AB028973 1.28
 hypothetical protein MGC10940 NM_018230 1.28
 hypothetical protein FLJ10814 AF217514 1.28
 lectin-cholesterol acyltransferase NM_018230 1.28
 three prime repair exonuclease 1 NM_000229 1.28
 carnitine palmitoyltransferase 1, muscle AJ243797 1.28
 U62733 1.28
 hypothetical protein U62733 1.28
 DNA-dependent protein kinase catalytic subunit-interacting protein 2 A1217472 1.28
 Consensus includes gb:AC004144 /DEF=Homo sapiens chromosome 19, overlapping cosmids R28707 and R34001 /FEA=CDS /DB_XREF=gi:2887496 /UG=Hs.307350 Homo sapiens cDNA: AC004144 1.28
 integrin, beta 5 BE138575 1.28
 transmembrane trafficking protein NM_006827 1.28
 RA-regulated nuclear matrix-associated protein NM_016448 1.28
 desrin (actin depolymerizing factor) NM_006870 1.28
 protein phosphatase 1, regulatory (inhibitor) subunit 15A U83961 1.28
 DNKZP434F195 protein BE222901 1.28
 hypothetical protein FLJ10101 NM_024718 1.28
 Consensus includes gb:AF123462 /DEF=Homo sapiens BAC526N18 neurexin III gene, partial cds /FEA=CDS /DB_XREF=gi:4240565 /UG=Hs.247837 Homo sapiens BAC526N18 neurexin III gene, partial cds: AF123462 1.28
 hypothetical protein MGC3133 NM_031287 1.28
 hypothetical protein FLJ20718 AV584285 1.28
 melanin 2 NM_006554 1.28

RAN binding protein 2	D47063	1,28
adenosine A2a receptor	NM_000675	1,28
thyroid hormone receptor-associated protein, 95-kD subunit	AA314406	1,28
KIAA0947 protein	BC004902	1,28
son of sevenless homolog 1 (Drosophila)	L13857	1,28
cell division cycle 27	AL566824	1,28
CGI-40 protein	AA150165	1,28
elastase 2, neutrophil	NM_001972	1,28
RAP1, GTPase activating protein 1	NM_002885	1,28
KRAB-zinc finger protein SZF1-1	NM_016089	1,28
protein disulfide isomerase related protein (calcium-binding protein, intestinal-related)	BC006344	1,28
hypothetical protein FLJ23109	NM_024814	1,28
mitochondrial ribosomal protein L20	NM_017971	1,28
pancreatic lipase	NM_000936	1,28
bone morphogenetic protein receptor, type II (serine/threonine kinase)	U20165	1,28
hypothetical protein FLJ22189	NM_024085	1,28
ribosomal protein L23a	U43701	1,28
polymerase (RNA) II (DNA directed) polypeptide C (33kD)	AJ224143	1,28
chromosome 1 open reading frame 16	BF509099	1,28
olfactory receptor, family 2, subfamily A, member 9 pseudogene	AA731709	1,28
ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2	X65909	1,28
phospholipid transfer protein	NM_006227	1,28
splicing factor protein/glutamine rich (polypyrimidine tract binding protein associated)	NM_005066	1,28
hypothetical protein MGC5528	NM_024094	1,28
splicing factor, arginine/serine-rich 4	NM_005626	1,28
hypothetical protein FLJ20345	NM_017777	1,28
antigen p87 (melanoma associated) identified by monoclonal antibodies 133.2 and 96.5	NM_005929	1,28
hypothetical protein MGC2835	NM_024072	1,28
ferredoxin-like protein 1 (farnesyl pyrophosphate synthetase, dimethylallyltransferase, geranyltransferase)	AL022163	1,28
protein phosphatase 1, regulatory subunit 7	AI060331	1,28
CDC5 cell division cycle 5-like (S. pombe)	NM_001253	1,28
TFP3 target gene 1	BC002709	1,28
EH-domain containing 1	AY007161	1,28
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsin, antitrypsin), member 2	NM_006220	1,28
BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)	NM_004725	1,28
muscle associated lymphoid tissue lymphoma translocation gene 1	NM_006785	1,27
RAB11A, member RAS oncogene family	AI215102	1,27
RNA binding motif, single stranded interacting protein 2	O28483	1,27
gb:NM_016104.1 /DEF=Homo sapiens PTD013 protein (PTD013), mRNA, /FEA=mRNA /GEN=PTD013 /PROD=PTD013 /PDB=Hs_279857 PTD013 protein /DB_XREF=gi:7706668 /UG=Hs_279857	NM_016104	1,27
zinc finger protein 282	AA521508	1,27
mitochondrial ribosomal protein S18A	AB049952	1,27
eukaryotic translation initiation factor 4 gamma, 2	NM_001418	1,27
hypothetical protein FLJ23053	NM_022907	1,27
myxovirus (influenza) resistance 1, homolog of murine (interferon-inducible protein p78)	NM_002462	1,27
baculoviral IAP repeat-containing 7 (livin)	NM_022161	1,27
sperm associated antigen 9	AK023512	1,27
ring finger protein 1	Z14000	1,27
hypothetical protein FLJ10587	NM_018149	1,27
cytosolic ovarian carcinoma antigen 1	AK000353	1,27
BCL2-associated athanogene 5	AA457021	1,27
artuin alien mating type information regulation 2 homolog 3 (S. cerevisiae)	AW280092	1,27
KIAA0776 protein	AK023092	1,27
DKF2P38D2223 protein	BE567813	1,27
ATPase, Cu++ transporting, alpha polypeptide (Menkes syndrome)	NM_006844	1,27
Irb (bacterial cardiolipase synthase)-like	AB041926	1,27
Mishapex/NIK-related kinase	AI373539	1,27
HIV TAT specific factor 1	AK023421	1,27
SEC7 homolog	AK023421	1,27
hypothetical protein	AW873594	1,27
Consensus includes gb:AW873564 /FEA=EST /DB_XREF=gi:8007617 /DB_XREF=eat:h62c10.x1 /CLONE=IMAGE:3041970 /UG=Hs_155485 huntingtin interacting protein 2	AW873564	1,27
COX15 homolog, cytochrome c oxidase assembly protein (yeast)	NM_004376	1,27
glutamine-fructose-6-phosphate transaminase 1	NM_002056	1,27
hypothetical protein FLJ21106	NM_025097	1,27

hypothetical protein FLJ10008 1.26
 small GTP-binding protein 1.26
 RNA 3'-terminal phosphate cyclase 1.26
 H1 histone family, member 2 1.26
 Consensus includes gb-BG178274 /FEA=EST /DB_XREF=gi:12684977 /DB_XREF=est:602330318f1 /CLONE=IMAGE:4431676 /UG=Hs.278895 postmeiotic segregation incrt BG178274 1.26
 Consensus includes gb-AA527409 /FEA=EST /DB_XREF=gi:2268568 /DB_XREF=est:ng4107.x1 /CLONE=IMAGE:937381 /UG=Hs.301667 Homo sapiens mRNA; cDNA DKF. AA527499 1.26
 heterogeneous nuclear ribonucleoprotein U (scarfold attachment factor A) 1.26
 Consensus includes gb-AL121777 /DEF=Human DNA sequence from clone RPS-1057D4 on chromosome 20 Contains a Spermidine synthase (SPDSY) pseudogene, a CpG is AL121777 1.26
 cullin 4A 1.26
 RNA (guanine-7-) methyltransferase 1.26
 arrestin, beta 1 1.26
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 1.26
 hypothetical protein DC30 1.26
 poly(A) binding protein, cytoplasmic 3 1.26
 hypothetical protein R33729.1 1.26
 GTPase regulator associated with the focal adhesion kinase pp125 1.26
 N-acetylglucosamine-phosphate mutase 1.26
 cutaneous T-cell lymphoma-associated tumor antigen sc20-4, differentially expressed nucleolar TGF-beta 1 target protein (DENT1) 1.26
 protocadherin beta 1 1.26
 hypothetical protein FLJ22405 1.26
 Consensus includes gb-AW409974 /FEA=EST /DB_XREF=gi:6935515 /DB_XREF=est:fn03d12.y1 /CLONE=IMAGE:2861094 /UG=Hs.332750 Homo sapiens cDNA FLJ10100 1.26
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a-like 1 1.26
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 5 1.26
 uncharacterized bone marrow protein BM029 1.26
 zinc finger protein 197 1.26
 KIAA1491 protein 1.26
 cytochrome c oxidase subunit VIc 1.26
 dynein light chain-A 1.26
 Consensus includes gb-AA310709 /FEA=EST /DB_XREF=gi:1963058 /DB_XREF=est:EST181521 /UG=Hs.249187 Homo sapiens BAC clone RP11-121A8 from 7p14-p13 1.26
 EH-domain containing 1 1.26
 apolipoprotein E 1.26
 gqg1 SNAP receptor complex member 2 1.26
 dual specificity phosphatase 12 1.26
 laminin receptor 1 (87kD, ribosomal protein SA) 1.26
 Td translation elongation factor, mitochondrial 1.26
 hypothetical protein FLJ13909 1.26
 NAD kinase 1.26
 chromosome 22 open reading frame 2 1.26
 hypothetical protein FLJ10709 1.26
 karyopherin (importin) beta 1 1.26
 actin cross-linking factor 1.26
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 11 (CHL1-like helicase homolog, S. cerevisiae) 1.26
 diaphenous homolog 1 (Drosophila) 1.26
 MAOS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C) 1.26
 calcium channel, voltage-dependent, gamma subunit 2 1.26
 Cbort10 protein 1.26
 nuclear receptor subfamily 6, group A, member 1 1.26
 flamin A, alpha (actin binding protein 280) 1.26
 hypothetical protein FLJ21272 1.26
 zinc finger RNA binding protein 1.26
 KIAA1411 protein 1.26
 hypothetical protein FLJ11749 1.26
 microsomal glutathione S-transferase 2 1.26
 DKF ZP547E2110 protein 1.26
 hypothetical protein FLJ10704 1.26
 ganglioside-induced differentiation-associated protein 1 1.26
 polycyochrome c synthase (cytochrome c heme-lyase) 1.26
 folypolyglutamate synthase 1.26
 Williams-Beuren syndrome chromosome region 1 1.26
 hypothetical protein FLJ13439 1.26
 Consensus includes gb-AI076186 /FEA=EST /DB_XREF=gi:3405364 /DB_XREF=est:oz01g01.x1 /CLONE=IMAGE:1674096 /UG=Hs.284275 Homo sapiens PAK2 mRNA, com 1.26
 beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase 1) 1.26

RAS p21 protein activator 2	NM_005506	1.26
karyopherin alpha 1 (Importin alpha 5)	BC002374	1.26
germ cell specific Y-box binding protein	NM_015982	1.26
nuclear cap binding protein subunit 1, 80kD	BG100193	1.26
eukaryotic translation initiation factor 3, subunit 6 (48kD)	BC000734	1.26
BTG family, member 2	BG339064	1.26
CD4-141 protein	NM_016072	1.26
secretory carrier membrane protein 1	AV745949	1.25
nuclear factor I/B	U70862	1.25
NLL sepin-like fusion	NM_006640	1.25
homo box B5	A052747	1.25
Consensus includes gb:AI017564 /FEA=EST /DB_XREF=gi:3231900 /DB_XREF=est:ou23e03.x1 /CLONE=IMAGE:1627132 /UG=Hs.95631 Human normal keratinocyte mRNA A017564	BG054916	1.25
patched homolog (Drosophila)	NM_004661	1.25
Consensus includes gb:AW519168 /FEA=EST /DB_XREF=gi:7157250 /DB_XREF=est:ha52a08.x1 /CLONE=IMAGE:2877302 /UG=Hs.163271 Human alpha-1 Ig germline C- α AW519168	AB002374	1.25
KIAA0378 protein	AF042162	1.25
cytochrome c oxidase subunit Va pseudogene 1	BE895285	1.25
golgi SNAP receptor complex member 2	NM_024668	1.25
hypothetical protein FLJ20288	NM_024668	1.25
Consensus includes gb:AW452796 /FEA=EST /DB_XREF=gi:5993572 /DB_XREF=est:UJ-H-B13-alk-4-06-0-UJ.s1 /CLONE=IMAGE:3069059 /UG=Hs.120769 Homo sapiens cDN AW452796	AW452796	1.25
hsp70-interacting protein	NM_012287	1.25
KIAA0033 protein	AU153138	1.25
KIAA0923 protein	NM_003122	1.25
serine protease inhibitor, Kazal type 1	NM_004661	1.25
CDC23 (cell division cycle 23, yeast, homolog)	AF798790	1.25
KIAA0728 protein	AL042733	1.25
BRCA1 associated protein	NM_017913	1.25
Consensus includes gb:AV691323 /FEA=EST /DB_XREF=gi:10293166 /DB_XREF=est:AV691323 /CLONE=GKCEWF11 /UG=Hs.20566 UDP glycosyltransferase 1 family, polyf AV691323	AV691323	1.25
putative nucleic acid binding protein RY-1	AW136988	1.25
hippocampin-like 1	X76302	1.25
KIAA0350 protein	BE917568	1.25
hypothetical protein FLJ13912	AA731693	1.25
glyoxalase 1	AI421812	1.25
erythropoietin	NM_006708	1.25
casein kinase 1, alpha 1	AF053356	1.25
fibronectin 1	AF119811	1.25
Hsp90-associating relative of Cdc37	AK026737	1.25
zinc finger protein 134 (clone pRZ-15)	NM_017913	1.25
PR domain containing 2, with ZNF domain	NM_003435	1.25
argininosuccinate lyase	AV724783	1.25
KIAA0828 protein	NM_000048	1.25
platelet derived growth factor C	AB020635	1.25
golgi autoantigen, golgin subfamily a, 5	NM_016205	1.25
RPR5-mediating protein	NM_005113	1.25
granulin	AW514900	1.25
endothelin receptor type B	AK023348	1.25
HLEMR homolog 7b	M74821	1.25
aminoacidpate-semialdehyde dehydrogenase-phosphopantetheinyl transferase	W93807	1.25
hypothetical protein FLJ14872	AF151057	1.25
solute carrier family 18 (monocarboxylic acid transporters), member 3	AA535065	1.25
cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila)	AA853175	1.25
WNT1 inducible signaling pathway protein 2	BC000059	1.25
small inducible cytokine subfamily B (Cys-X-Cys), member 11	NM_003881	1.25
chromogranin A (parathyroid secretory protein 1)	AF002585	1.25
CLIP-associating protein 1	NM_001275	1.25
hypothetical protein DKFZp547M136 similar to widely-interspaced zinc finger motifs	AA176798	1.25
Consensus includes gb:AW449390 /FEA=EST /DB_XREF=gi:6990166 /DB_XREF=est:UJ-H-B13-aki-g-10-0-UJ.s1 /CLONE=IMAGE:2734650 /UG=Hs.257150 ESTs, Moderately AW449390	AI828531	1.25
signal transducer and activator of transcription 6, interleukin-4 induced	BC004973	1.25
umilto homolog 1 (Drosophila)	NM_014676	1.25
CDC-like kinase 3	NM_003992	1.25
polycystic kidney disease 1 (autosomal dominant)	NM_000296	1.25
TLA1 cytosolic granule-associated RNA binding protein-like 1	NM_003252	1.25
isoprenylcysteine carboxyl methyltransferase	NM_012405	1.25
ABO blood group (transferase A, alpha 1,3-N-acetylgalactosaminyltransferase; transferase B, alpha 1,3-galactosyltransferase)	NM_020469	1.25

A20-binding inhibitor of NF-kappaB activation-2
 Consensus includes gb:269637.1 /DEF=H.sapiens mRNA sequence (16p11.2). /FEA=mRNA /DB_XREF=gi:29573 /UG=Hs.307208 H.sapiens mRNA sequence (16p11.2)
 uncharacterized hematopoietic stem/progenitor cells protein MDS033
 Rho guanine exchange factor (GEF) 11
 Consensus includes gb:A852772 /FEA=EST /DB_XREF=gi:5745082 /DB_XREF=est:ws6005.x1 /CLONE=IMAGE:2491113 /UG=Hs.300865 Homo sapiens clone BCSynL38
 transmembrane 4 superfamily member 6
 hypothetical protein FLJ20030
 growth arrest and DNA-damage-inducible, alpha
 HSPC133 protein
 hypothetical protein FLJ22347
 luciferin transferase 2 (secretor status included)
 diacylglycerol kinase, beta
 uncharacterized hypothalamus protein HT007
 protein phosphatase 3 (formery 2B), catalytic subunit, beta isoform (calcineurin A beta)
 BCL2adenovirus E1B 19kD interacting protein 1
 forhead box H1
 ATP-binding cassette, sub-family G (WHITE), member 1
 DKFZP368B1621 protein
 HSPC134 protein
 carboxypeptidase D
 hypothetical protein FLJ11273
 x 005 protein
 chromodomain helicase DNA binding protein 4
 hypothetical protein FLJ14390
 myeloid lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10
 spinlin
 zinc finger protein 108
 branched chain alpha-ketoadid dehydrogenase kinase
 cell division cycle 2-like 1 (PTSLRE proteins)
 hypothetical protein FLJ13910
 phospholipase C, gamma 2 (phosphatidylinositol-specific)
 carboxypeptidase Z
 hypothetical protein BC022942
 Consensus includes gb:AL390153.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762G113 (from clone DKFZp762G113). /FEA=mRNA /DB_XREF=gi:5368841 /UG=Hs.208339
 hypothetical protein FLJ20581
 baculoviral AP repeat-containing 1
 candidate mediator of the p53-dependent G2 arrest
 putative DNA binding protein
 MCM6 minichromosome maintenance deficient 6 (MISS homolog, S. pombe) (S. cerevisiae)
 ARP1 actin-related protein 1 homolog B, centractin beta (yeast)
 nasopharyngeal epithelium specific protein 1
 KIAA0354 protein
 heterogenous nuclear ribonucleoprotein H1 (H)
 hypothetical protein FLJ20548
 retinoic acid induced 16
 beta-2-microglobulin
 proteasome (prosome, macropain) 26S subunit, non-ATPase, 11
 seductin 3 (gamma)
 polymerase (DNA directed), mu
 KIAA1098 protein
 dynamin 1
 copin VI (neuronal)
 hypothetical protein FLJ10208
 hypothetical protein MGC3282
 hypothetical protein FLJ10743
 suppressor of potassium transport defect 3
 tumor differentially expressed 1
 Consensus includes gb:S79287.1 /DEF=CD4 receptor (exons 1 and 2) human, T-lymphocyte, mRNA, 3429 nt. /FEA=mRNA /GEN=CD4 receptor /PROD=CD4 receptor /DB_XS_S79287
 Splicing factor, arginine/serine-rich, 46kD
 hypothetical protein FLJ22035
 pancreatic polypeptide
 Consensus includes gb:AW128021 /FEA=EST /DB_XREF=gi:8116965 /DB_XREF=est:xe93h03.x1 /CLONE=IMAGE:2615477 /UG=Hs.243537 ESTs, Highly similar to AC00504
 JMS protein
 BC000464

KIAA0993 protein	AL536319	1,24
hypothetical protein FLJ10379	NM_018079	1,24
ryliokininase (H. influenzae) homolog	NM_005108	1,24
tubulin beta-5	BC002654	1,24
hypothetical gene MGC9753	AI703342	1,24
cystathione conjugate-beta lyase, cytoplasmic (glutamine transaminase K, kynurenine aminotransferase)	NM_004059	1,24
signal transducing adaptor molecule (SH3 domain and ITAM motif) 1	NM_003473	1,24
ribosomal protein S8	A6976278	1,24
BCL2-associated X protein	NM_004324	1,24
COX15 homolog, cytochrome c oxidase assembly protein (yeast)	BC002382	1,24
hypothetical protein HSPC177	NM_018410	1,24
putative-rich element binding protein A	NM_005859	1,24
centrosomal protein 2	AF022655	1,24
paired box gene 2	NM_003988	1,24
hypothetical protein dJ12208.2	NM_020466	1,24
Cas-B1-M (murine) ectropic retroviral transforming sequence c	NM_012116	1,24
Ca2+-promoted Ras inactivator	NM_006989	1,24
Consensus includes gb:AK000834.1 /DEF=Homo sapiens cDNA FLJ20827 flt. clone ADKA03543. /FEA=mrna /DB_XREF=gi:7021152 /UG=Hs.306409 Homo sapiens cDNA	AK000834	1,24
solute carrier family 7 (cationic amino acid transporter, y+ system), member 8	NM_012244	1,24
fatty acid desaturase 1	BE540552	1,24
hypothetical protein FLJ20327	NM_017787	1,24
Consensus includes gb:AL035588 /DEF=Human DNA sequence from clone 696P19 on chromosome 6p12.3-21.2. Contains the gene for TFEB, an NPM1 (Nucleophosmin, Nur	AL035588	1,24
hypothetical protein FLJ10385	NM_018081	1,24
homeo box B5	NM_002147	1,24
LRP16 protein	NM_014067	1,24
KIAA0102 gene product	NM_014752	1,24
FX10 domain-containing ion transport regulator 1 (phospholemman)	NM_005031	1,24
Consensus includes gb:AL583123 /FEA=EST /DB_XREF=gi:12951784 /DB_XREF=est:AL583123 /CLONE=CS0DL009YN08 (3 prime) /UG=Hs.294005 Homo sapiens, clone 1h	AL583123	1,24
hypothetical protein FLJ13220	NM_021927	1,24
topoisomerase (DNA) II alpha (170kD)	AU159942	1,24
protocadherin gamma subfamily A, 10	AF152503	1,24
polymerase (DNA directed), epsilon 2	NM_002692	1,24
eukaryotic translation initiation factor 3, subunit 3 (gamma, 40kD)	NM_003756	1,24
microtubule-associated protein 4	M64571	1,24
zinc finger protein 145 (Kruppel-like, expressed in promyelocytic leukemia)	NM_006006	1,24
ring finger protein 14	AB022663	1,24
COI-42 protein	AF167438	1,24
annexin A2 pseudogene 2	M62898	1,24
Consensus includes gb:BF448531 /FEA=EST /DB_XREF=gi:11514556 /DB_XREF=est:17n72e02.x1 /CLONE=IMAGE 3570146 /UG=Hs.248979 ESTs, Moderately similar to ALI	BF448531	1,24
ribosomal protein L22	D17652	1,24
hypothetical protein FLJ11159	NM_018343	1,24
ARF3 actin-related protein 3 homolog (yeast)	AL544094	1,24
KIAA0116 protein	AL353587	1,24
Huntingtin interacting protein K	AL581473	1,24
DKFZP434B103 protein	NM_016400	1,24
KIAA0155 gene product	AF078842	1,24
SRF (sex determining region Y)-box 11	NM_014633	1,24
diacylglycerol O-acyltransferase homolog 1 (mouse)	AB028641	1,24
adenomatous polyposis coli	NM_012079	1,24
ring finger protein 24	M74088	1,24
cylindromatosis (turban tumor syndrome)	NM_007219	1,24
chromosome 11 open reading frame 13	AK024212	1,24
thrombopoietin (myeloproliferative leukemia virus oncogene ligand, megakaryocyte growth and development factor)	M91083	1,24
choline kinase	D32047	1,24
hypothetical protein FLJ20258	AIS91328	1,24
Consensus includes gb:A0807507 /FEA=EST /DB_XREF=gi:5394073 /DB_XREF=est:hw48e10.x1 /CLONE=IMAGE 2358858 /UG=Hs.126496 Homo sapiens chromosome 19, c	NM_017729	1,24
TRAM-like protein	NM_012288	1,24
protein phosphatase 1, catalytic subunit, gamma isoform	NM_002710	1,24
nucleoporin 50KD	NM_007172	1,24
zinc finger protein, subfamily 1A, 1 (Ikaros)	NM_006060	1,24
KIAA0552 gene product	NM_014731	1,24
synaptic vesicle protein 28 homolog	NM_014848	1,24

synovial sarcoma, X breakpoint1 2	BC002818	1.17
fibroblast growth factor 8	NM_020986	1.17
opoid binding protein/cell adhesion molecule-like	NM_002545	1.17
G protein-coupled receptor 44	AF118265	1.17
Fzr1 protein	AB013463	1.17
guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 2	NM_002070	1.17
putative 28 kDa protein	NM_020143	1.17
like-glycosyltransferase	AB011181	1.17
lanthipeptin (importin) beta 2	NM_002270	1.17
cell division cycle 42 (GTP binding protein, 25kD)	R37664	1.17
SCO cytochrome oxidase deficient homolog 2 (yeast)	NM_005138	1.17
jagged 2	AF029778	1.17
v-rat-1 murine leukemia viral oncogene homolog 1	X06409	1.17
frizzled homolog 5 (Drosophila)	NM_003468	1.17
Consensus includes gb:H24302 /FEA=EST /DB_XREF=gi:892997 /DB_XREF=est:ym51e07.s1 /CLONE=IMAGE:51738 /UG=Hs.23127 ESTs	H24302	1.17
hypothetical protein MGC4276 similar to CG8188	BC002875	1.17
Consensus includes gb:A1188739 /FEA=EST /DB_XREF=gi:3737377 /DB_XREF=est:q679c01.x1 /CLONE=IMAGE:1745184 /UG=Hs.326416 Homo sapiens mRNA; cDNA DKF	A1188739	1.17
MYC-associated zinc finger protein (putative-binding transcription factor)	A471665	1.17
hypothetical protein FLJ20657	NM_017921	1.17
splicing factor, arginine/serine-rich 2, interacting protein	A954932	1.17
hypothetical protein FLJ12448	NM_022895	1.17
G antigen, family B, 1 (prostate associated)	NM_003785	1.17
GDP dissociation inhibitor 1	NM_001493	1.17
Consensus includes gb:AL137784 /DEF=Human DNA sequence from clone RP1-198J3 on chromosome 6 Contains ESTs, STSs, GSSs and CpG islands. Contains part of the	AL137784	1.17
Consensus includes gb:X81636.1 /DEF=H.sapiens clathrin light chain a gene. /FEA=mRNA /DB_XREF=gi:704480 /UG=Hs.295689 H.sapiens clathrin light chain a gene	X81636	1.17
Ras-GTPase activating protein SH3 domain-binding protein 2	AB014560	1.17
CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, E118, E130, EL32 and G344)	BF983379	1.17
hypothetical protein, similar to (U06944) PRAJA1	AL530462	1.17
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 2	NM_002575	1.17
melanoma antigen, family D, 1	AF217963	1.17
endosome-associated FYVE-domain protein	NM_014723	1.17
mucin and cadherin-like	NM_021924	1.17
ROD1 regulator of differentiation 1 (S. pombe)	AW190873	1.17
KIAA1696 protein	NM_016621	1.17
ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeier-Vogt disease)	AF077973	1.17
G protein-coupled receptor 51	AF095723	1.17
alko-keto reductase family 1, member C3 (3-alpha hydroxysteroid dehydrogenase, type II)	AB018580	1.17
leucocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2	U82276	1.17
origin recognition complex, subunit 4-like (yeast)	NM_002552	1.17
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14)	NM_002490	1.17
mitofusin 1	NM_017827	1.17
hypothetical protein FLJ10545	NM_018132	1.17
integral membrane protein 1	NM_002219	1.17
G protein-coupled receptor 56	NM_005682	1.17
JAK binding protein	AA877218	1.17
stem cell growth factor: lymphocyte secreted C-type lectin	NM_002975	1.17
heterogeneous nuclear ribonucleoprotein A3	NM_005758	1.17
G protein-coupled receptor 4	U35399	1.17
mitochondrial ribosomal protein S31	NM_005830	1.17
Consensus includes gb:AF189251.1 /DEF=Homo sapiens cytomegalovirus partial fusion receptor mRNA, partial cds. /FEA=mRNA /PROD=cytomegalovirus partial fusion recat	AF189251	1.17
SAC2 suppressor of actin mutations 2-like (yeast)	AL390171	1.17
KIAA0751 gene product	AF007156	1.17
protein phosphatase 2 (formerly 2A), regulatory subunit B' (PR 72), alpha isoform and (PR 130), beta isoform	NM_002718	1.17
polymerase (RNA) II (DNA directed) polypeptide K (7.0kD)	AL558030	1.17
S-adenosylhomocysteine hydrolase-like 1	NM_006621	1.17
excision repair cross-complementing rodent repair deficiency, complementation group 3 (xeroderma pigmentosum group B complementing)	NM_000122	1.17
dihydropyrimidine branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease)	M27093	1.17
calcium binding atopy-related autoantigen 1	AK022697	1.17
phenylalanine-RNA synthetase	NM_006567	1.17
hypothetical protein MGC4808	NM_024516	1.17
SHB7 suppressor of RNA polymerase B homolog (yeast)	U46837	1.17
3-hydroxyethyl-3-methylglutaryl-Coenzyme A lyase (hydroxymethylglutarylaciduria)	NM_000191	1.17
gastric inhibitory polypeptide	NM_004123	1.17

hypothetical SBBI03 protein	AL583171	1.17
ceroid-lipofuscinosis, neuronal 5	A0911687	1.17
hypothetical protein MGC10765	NM_024345	1.17
Rho guanine nucleotide exchange factor (GEF) 5	NM_005435	1.17
X-ray repair complementing defective repair in Chinese hamster cells 3	AK022829	1.17
KIAA1155 protein	AB032981	1.17
similar to yeast BET3 (S. cerevisiae)	NM_014408	1.17
mitogen-activated protein kinase kinase 5	NM_002757	1.17
DKFZP772M231 protein	AL117480	1.17
similar to yeast Upt3, variant A	AI684141	1.17
heart shock 70kD protein 4	NM_023011	1.17
cannabinoid receptor 1 (brain)	AB023420	1.17
gpJF308302.1 IDEF-Homo sapiens serologically defined breast cancer antigen NY-BR-98 mRNA, complete cds. /FEA=mRNA	NM_016083	1.17
protease, cysteine, 1 (legumain)	AF308302	1.17
type 1 tumor necrosis factor receptor shedding aminopeptidase regulator	AW450751	1.17
histone deacetylase 5	D55896	1.17
ring finger protein 27	AF106037	1.17
tumor endothelial marker 1 precursor	NM_005474	1.17
collagen, type IV, alpha 3 (Goodpasture antigen) binding protein	NM_030912	1.17
mitogen-activated protein kinase-activated protein kinase 2	NM_020404	1.17
phosphate cytidylyltransferase 2, ethanolamine	NM_005713	1.17
Down syndrome critical region gene 5	AI141802	1.17
activated p21cdc42Hs kinase	BC000351	1.17
protein tyrosine kinase 9-like (A6-related protein)	AB035745	1.17
hypothetical protein FLJ13164	NM_007284	1.17
solute carrier family 15 (H+/peptide transporter), member 2	NM_022776	1.16
regulator of nonsense transcripts 2	NM_021082	1.16
DKFZP564A043 protein	BE350145	1.16
mucin 8, tracheobronchial	NM_015342	1.16
zinc finger protein 148 (pH2.52)	U14383	1.16
splicing factor, arginine/serine-rich 9	L04282	1.16
numa homolog (Drosophila)	NM_003769	1.16
PBX/Notch1 1 homeobox 1	NM_003744	1.16
peroxisome receptor 1	NM_004571	1.16
PTK6 protein tyrosine kinase 6	AW468717	1.16
sorcin	NM_005975	1.16
adaptor-related protein complex 3, beta 1 subunit	L12387	1.16
hypothetical protein FLJ10781	AW058575	1.16
protein phosphatase 8, catalytic subunit	NM_018215	1.16
shvelar soft part sarcoma chromosome region, candidate 1	NM_018294	1.16
zinc finger protein 95 homolog (mouse)	NM_024083	1.16
XIAP associated factor-1	NM_014589	1.16
cytochrome P450, subfamily XXIV (vitamin D 24-hydroxylase)	NM_017523	1.16
general transcription factor II, 1	AI432196	1.16
zinc finger protein, subfamily 1A, 3 (Atf1os)	NM_000782	1.16
nuclear receptor co-repressor 1	AF035737	1.16
glycine cleavage system protein H (aminomethyl carrier)	NM_012481	1.16
transforming growth factor, beta 2	NM_006311	1.16
Conserved gene telomeric to alpha globin cluster	AW237404	1.16
glucosamine (N-acetyl)-6-sulfatase (Sanfilippo disease IIID)	NM_003238	1.16
TNF receptor-associated factor 2	BC004185	1.16
hypothetical protein FLJ22893	NM_002076	1.16
Kruppel-like factor 15	NM_022750	1.16
pleomorphic adenoma gene-like 1	NM_014079	1.16
platelet-activating factor acetylhydrolase, isoform 1b, alpha subunit (45kD)	NM_006718	1.16
glucosamine-6-phosphatase isomerase	L13386	1.16
KIAA0852 protein	NM_005471	1.16
zinc finger protein 140 (clone pHZ-39)	AC004542	1.16
KIAA0152 gene product	NM_003440	1.16
	BC000371	1.16

protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform
homo box D5
DnaJ (Hsp40) homolog, subfamily B, member 6
Kruppel-like factor 3 (intestinal)
fragile X mental retardation 1
cyclin L, enla-6a
karyopherin (importin) beta 2
UDP-N-acetylglucosamine pyrophosphorylase 1
cell division cycle 25B
FtsJ homolog 2 (E. coli)
non-metastatic cells 3, protein expressed in
progesterone receptor membrane component 1
Mouse Mammary Tumor Virus Receptor homolog
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2
ubiquitin specific protease 15
chloride intracellular channel 1
KIAA0871 gene product
hypothetical protein MGC955
cell cycle progression 2 protein
olfactory receptor, family 2, subfamily F, member 1
epimorphin
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform
gb:NM_018611.1 /DEF=Homo sapiens hypothetical protein PRO1966 (PRO1966), mRNA, /FEA=mRNA /GEN=PRO1966 /PROD=hypothetical protein PRO1966 /DB_XREF=gi:10203952 /DB_XREF=est:u4307.x1 /CLONE=IMAGE:2804508 /UG=Hs.288760 Homo sapiens cDNA FLJ12327, AW438464
cyclin M3
KIAA0560 gene product
hypothetical protein FLJ11786
signal peptidase complex (18kD)
potassium large conductance calcium-activated channel, subfamily M beta member 3
sodium channel, nonvoltage-gated 1, gamma
Hiv-1 Tat interactive protein, 60 kDa
sarcosine dehydrogenase
tumor necrosis factor receptor superfamily, member 6b, decoy
SPTF-associated factor 65 gamma
platelet-derived growth factor alpha polypeptide
breast cancer 1, early onset
karyopherin (importin) beta 1
microtubule-associated protein, RP/EB family, member 2
tumor protein D52
hypothetical protein FLJ13611
KIAA0157 protein
adaptor-related protein complex 1, sigma 1 subunit
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ceroid-f lipofuscinosis, neuronal 2, late infantile (Jansky-Bielschowsky disease)
Isovaleryl Coenzyme A dehydrogenase
zinc finger protein 215
adducin 3 (gamma)
kinesin-like 2
neuropilin 2 Y receptor Y1
immunoglobulin (CD79A) binding protein 1
neuritin 1
XPM1C2 protein
signaling lymphocytic activation molecule
acyl-Coenzyme A oxidase 2, branched chain
arachidonate 5-lipoxygenase
transport-secretion protein 2.2,
ethanolamine kinase
Consensus includes gb:AW438464 /FEA=EST /DB_XREF=gi:6973770 /DB_XREF=est:u4307.x1 /CLONE=IMAGE:2804508 /UG=Hs.288760 Homo sapiens cDNA FLJ12327, AW438464
gb:M52596.1 /DEF=Homo sapiens immunoglobulin kappa-chain VK-1 (IgK) mRNA, complete cds, /FEA=mRNA /GEN=IgK /PROD=immunoglobulin kappa-chain VK-1 /DB_XRI M52596
arachidonate 12-lipoxygenase pseudogene 2
complement component 4B
cyclin-dependent kinase (CDC2-like) 10
hypothetical protein FLJ11101
glucocorticoid modulatory element binding protein 2

AF163473 1.16
A432470 1.16
AF080569 1.16
AF132818 1.16
A1743037 1.16
NM_020307 1.16
A1653355 1.16
S73498 1.16
NM_021873 1.16
NM_013393 1.16
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AF064842 1.16
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NM_003037 1.16
NM_003500 1.16
A1361850 1.16
AF055000 1.16
AL137750 1.16
AW438464 1.16
AF020774 1.16
AL048547 1.16
AF153430 1.16
NM_018322 1.16
AA045183 1.16

NIMA (never in mitosis gene a)-related kinase 1	AI938517	1.16
tripetide motif-containing 10	X90539	1.16
tripetide motif-containing 2	AW071795	1.16
double homeobox, 2	NM_012147	1.16
CD209 antigen	AF280866	1.16
phospholipase A2, group IVB (cytosolic)	NM_005090	1.16
hypothetical protein FLJ21877	NM_022065	1.16
phosphatidylcholine transfer protein	NM_021213	1.16
KIAA0252 protein	D87440	1.16
Consensus includes gb:AK021738.1 /DEF=Homo sapiens cDNA FLJ11876 fls. clone HEMBA1004752, highly similar to Homo sapiens mRNA for LAX-4p. /FEA=mRNA DB_XF AK021738		1.16
Consensus includes gb:Y15724 /DEF=Homo sapiens SERCA3 gene, exons 1-7 (and joined CDS) /FEA=mRNA DB_XREF=gi:3021395 /UG=Hs.5541 ATPase, Ca++ transport Y15724		1.16
unchacterized bone marrow protein BM046	NM_018460	1.16
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)	BE787213	1.16
protein kinase C, etc	NM_006255	1.16
inter-alpha (globulin) inhibitor, H2 polypeptide	NM_002216	1.16
KIAA1719 protein	AF052177	1.16
hypothetical protein FLJ23338	NM_024658	1.16
heterogeneous nuclear ribonucleoprotein H1 (H)	NM_024658	1.16
ribulose-5-phosphate-3-epimerase	BF983408	1.16
hypothetical protein DKFZp434L0117	BE984473	1.16
glycosyltransferase	NM_022778	1.16
microtubular-associated protein 4	NM_031302	1.16
Consensus includes gb:AF007131.1 /DEF=Homo sapiens clone 23963 mRNA sequence /FEA=mRNA DB_XREF=gi:2852805 /UG=Hs.48483 Homo sapiens clone 23963 mR	R72286	1.16
mitogen-activated protein kinase kinase kinase 1	AA744529	1.16
5-hydroxytryptamine (serotonin) receptor 5A	NM_024012	1.16
CD72 antigen	AF283777	1.16
dual specificity phosphatase 5	U16996	1.16
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fibroblast growth factor 12B	NM_004113	1.16
calreticulin	BE251303	1.16
ribosomal protein S6 kinase, 90kD, polypeptide 1	NM_002853	1.16
LIM domain kinase 2	NM_005569	1.16
Consensus includes gb:Y08772 /DEF=H. sapiens PABII pseudogene /FEA=CDS DB_XREF=gi:1770485 /UG=Hs.247983 H. sapiens PABII pseudogene	Y08772	1.16
fer-1-like 3, myoferlin (C. elegans)	AF207990	1.16
amyloid beta precursor protein (cytoplasmic tail) binding protein 2	AV681579	1.16
related RAS viral (v-ras) oncogene homolog, 2	A431843	1.16
Consensus includes gb:AK028867.1 /DEF=Homo sapiens cDNA: FLJ23214 fls. clone ADSU01585. /FEA=mRNA DB_XREF=gi:10439827 /UG=Hs.306891 Homo sapiens cDNA: AK028867	AK028867	1.16
inositol 1,4,5-trisphosphate receptor, type 3	NM_002224	1.16
myelin protein zero-like 1	AF095727	1.16
potassium voltage-gated channel, shaker-related subfamily, beta member 2	AF044253	1.16
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 4	NM_016129	1.16
TXK tyrosine kinase	NM_003328	1.16
epidermal growth factor receptor pathway substrate 15	BF213575	1.16
hypothetical protein R31240.1	A435954	1.16
DEADH (Asp-Glu-Ala-Asp-His) box polypeptide 11 (CHL 1-like helicase homolog, S. cerevisiae)	NM_004399	1.16
phosphatidylinositol-specific phospholipase D1	L11702	1.16
immunoglobulin heavy constant gamma 3 (G3m marker)	NM_025072	1.16
hypothetical protein FLJ14038	X95746	1.16
potassium channel, subfamily K, member 10 (TREK-2)	AI765051	1.16
PC4 and SFRS1 interacting protein 2	NM_021161	1.16
similar to KIAA0010 gene product (H. sapiens).	AF083020	1.16
hypothetical protein FLJ12517	AL096740	1.16
hypothetical protein FLJ10283	NM_023007	1.15
caspase 3, apoptosis-related cysteine protease	NM_018046	1.15
matrix metalloproteinase 19	NM_004346	1.15
KIAA1054 protein	U38321	1.15
calmodulin 2 (phosphorylase kinase, delta)	AB028977	1.15
protein tyrosine kinase 2 beta	NM_001743	1.15
liver cell autoantigen 1 (69kD)	U43522	1.15
enigma (LIM domain protein)	W67992	1.15
steroid regulatory element binding transcription factor 1	AA086229	1.15
multifunctional polypeptide similar to SAICAR synthetase and AIR carboxylase	NM_004176	1.15
	AA026852	1.15

kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)
 glutathione peroxidase 4 (phospholipid hydroperoxidase)
 chromatin assembly factor 1, subunit B (p60)
 Custer Ind. AW025284: wu95h.10.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:990806 /clone_end=3 /gb=AW025284 /gi=5878814 /ug=Hs.237946 /len=454
 hemochromatosis
 nuclear receptor subfamily 6, group A, member 1
 KIAA0878 protein
 vesicle-associated membrane protein 2 (synaptobrevin 2)
 hypothetical protein MGC3079
 hypothetical protein FLJ10074
 protein phosphatase 3 (former 2B), catalytic subunit, alpha isoform (calcineurin A alpha)
 p30 DBC protein
 zinc finger protein 144 (Mei-18)
 BTG family, member 2
 cytochrome P450, subfamily XB (steroid 11-beta-hydroxylase), polypeptide 1
 bone morphogenetic protein 1
 rotarotin 2
 proteoglycan 4, (megakaryocyte stimulating factor, articular superficial zone protein, camptodactyl, arthropathy, coxa vara, pericarditis syndrome)
 quiescin Q6
 protein lysine phosphatase-like (proline instead of catalytic arginine), member a
 olfactory receptor, family 2, subfamily B, member 6
 Consensus includes gb:BE46525 /FEA=EST /DB_XREF=gi:9512223 /DB_XREF=est:th94b10.x1 /CLONE=IMAGE:3195451 /UG=Hs.234773 Homo sapiens cDNA: FLJ22281
 KIAA1859 protein
 gamma-glutamyltransferase 1
 mitochondrial ribosomal protein S33
 desmoglein 3 (pemphigus vulgaris antigen)
 hypothetical protein FLJ2247.1
 huntingtin interacting protein 1
 acylphosphatase 2, muscle type
 KIAA0899 protein
 hypothetical protein FLJ20539
 hypothetical protein PRO0899
 VAMP (vesicle-associated membrane protein)-associated protein B and C
 eis variant gene 5 (eis-related molecule)
 core-binding factor, beta subunit
 hypothetical protein FLJ20398
 cyclin-dependent kinase 6
 Agt, S. cerevisiae, homolog of
 phosphodiesterase 4C, cAMP-specific (phosphodiesterase E1 dunce homolog, Drosophila)
 carbonic anhydrase VB, mitochondrial
 retinoid X receptor, beta
 testis zinc finger protein
 solute carrier family 21 (organic anion transporter), member 9
 Consensus includes gb:AK025097.1 /DEF=Homo sapiens cDNA: FLJ21444 /lis, clone COL04437, /FEA=mrna /DB_XREF=gi:10437544 /UG=Hs.306764 Homo sapiens cDNA: NM_001652
 aquaporin 6, kidney specific
 FAST thase
 gp25L2 protein
 cysteine and glycine-rich protein 1
 MCT-1 protein
 protein lysine phosphatase, non-receptor type 18 (brain-derived)
 receptor-interacting serine-threonine kinase 2
 hypothetical protein PRO1580
 solute carrier family 2 (facilitated glucose transporter), member 4
 protective protein for beta-galactosidase (galactosialidosis)
 hypothetical protein FLJ21799
 KIAA0435 gene product
 retinoic acid repressible protein
 homolog of mouse quaking OKI (KH domain RNA binding protein)
 degenerative spermatocyte homolog, lipid desaturase (Drosophila)
 protein phosphatase 4 regulatory subunit 2
 cytochrome b5 reductase 1 (BSR-1)
 Consensus includes gb:W74494 /FEA=EST /DB_XREF=gi:1384781 /DB_XREF=est:z475f10.s1 /CLONE=IMAGE:346507 /UG=Hs.284275 Homo sapiens PAK2 mRNA, complete W74494
 Consensus includes gb:AW500220 /FEA=EST /DB_XREF=gi:7112628 /DB_XREF=est:U1-HF-BN0-akd-c02-0-U1.r1 /CLONE=IMAGE:3076610 /UG=Hs.332496 Human DNA ss AW500220

cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila)
hypothetical protein FLJ14547 AL031588 1.15
G protein-coupled receptor kinase 6 AL049319 1.15
Consensus includes gb:AA591879 /FEA=EST /DB_XREF=gi:2380557 /DB_XREF=est:nn43g06.s1 /CLONE=IMAGE:1086682 /UG=Hs.66191 Homo sapiens clone 24875 mRNA /AA581879 1.15
Consensus includes gb:AK021890.1 /DEF=Homo sapiens cDNA FLJ11628 flg. clone HEMBA1004238. /FEA=mRNA /DB_XREF=gi:10432926 /UG=Hs.306610 Homo sapiens c AK021890 1.15
gb:NM_030807.1 /DEF=Homo sapiens glucose transporter protein 10 (GLUT10). mRNA. /FEA=mRNA /GEN=GLUT10 /PROD-glucose transporter protein 10 /DB_XREF=gi:13. NM_030807 1.15
hypothetical protein FLJ23584 NM_024588 1.15
hypothetical protein FLJ13111 NM_025082 1.15
phosphodiesterase 3B, cGMP-inhibited NM_000592 1.15
synovial sarcoma, X breakpoint 2 AF343880 1.15
Consensus includes gb:AF035594.1 /DEF=Homo sapiens protein kinase C-alpha mRNA, partial 3 UTR. /FEA=mRNA /DB_XREF=gi:3168857 /UG=Hs.279856 Homo sapiens p AF035594 1.15
mazlin 1, cartilage matrix protein NM_002379 1.15
male-enhanced antigen NM_014623 1.15
Fancconi anemia, complementation group G NM_004629 1.15
fatty-acid-Coenzyme A ligase, long-chain 3 AL525788 1.15
src family associated phosphoprotein 2 NM_003930 1.15
cone-binding factor, runt domain, alpha subunit 2; translocated to, 3 NM_005187 1.15
keratohelin alpha 3 (importin alpha 4) AF034756 1.15
KIAA0210 gene product A654161 1.15
Consensus includes gb:AL034429 /DEF=Human DNA sequence from clone RPS-894K16 on chromosome 20q11.3.11.13.13. Contains a COX6C (cytochrome c oxidase subunit) AL034429 1.15
pituitary tumor-transforming 3 NM_021000 1.15
chromosome 11 open reading frame 24 AA085185 1.15
Spogren syndrome antigen B (autoantigen La) NM_003142 1.15
hypothetical protein FLJ10787 NM_018209 1.15
gamma-glutamyltransferase 1 NM_013421 1.15
hypothetical protein FLJ12488 NM_031218 1.15
splicing factor, arginine/serine-rich 8 (suppressor-of-white-apricot homolog, Drosophila) AL023864 1.15
pregnancy-associated interferon L25664 1.15
hypothetical protein FLJ20045 NM_017638 1.15
hypothetical protein DKFZP547F082 AL512726 1.15
zinc finger protein 200 NM_003454 1.15
AND-1 protein AW772140 1.15
phosphatidylserine decarboxylase AL096768 1.15
DEAD(H)(Asp-Glu-Ala-Asp-His) box polypeptide 11 (CHL 1-like helicase homolog, S. cerevisiae) NM_030653 1.15
engulfment and cell motility 2 (ced-12 homolog, C. elegans) NM_022086 1.15
homo box B7 NM_004502 1.15
lymphoid enhancer-binding factor 1 AF268571 1.15
guanine DNA/chromatin binding motif W02593 1.15
Consensus includes gb:AL522296 /FEA=EST /DB_XREF=gi:12785789 /DB_XREF=est:AL522296 /CLONE=CSDB008YP08 (5 prime) /UG=Hs.17481 Homo sapiens clone 248 AL522296 1.15
SHC (Src homology 2 domain-containing) transforming protein 1 A089867 1.15
hypothetical protein FLJ10618 A1694452 1.15
tumor suppressing substrate transferable candidate 4 NM_005706 1.15
transporter 2, ATP-binding cassette, sub-family B (MDR/TAP) NM_000544 1.15
ELAV (embryonic lethal, abnormal vision, Drosophila)-like 1 (Hu antigen R) BC003376 1.15
hypothetical protein FLJ20260 AF323729 1.15
gb:M17565.1 /DEF=Human MHC class II DQ-beta associated with DRw6, DQw1 protein, complete cds. /FEA=mRNA /GEN=HLA-DQB1 /DB_XREF=gi:188188 /FL=gb:M17565 M17565 1.15
neuroblastoma-amplified protein NM_015909 1.15
hypothetical protein FLJ13855 NM_023079 1.15
keratohelin (importin) beta 2 A1307739 1.15
thyroid receptor interacting protein 15 A4496247 1.15
neuralin NM_005386 1.15
NIMA (never in mitosis gene a)-related kinase 3 A1191920 1.15
Consensus includes gb:AK024289.1 /DEF=Homo sapiens cDNA FLJ14207 fls. clone NT2RP3003185, weakly similar to TROPOMYOSIN 1, FUSION PROTEIN 33. /FEA=mRNA AK024289 1.15
prostaglandin 12 (prostaglandin) synthase NM_000961 1.15
EphA1 Z27403 1.14
G protein-coupled receptor 65 NM_003608 1.14
KIAA0117 protein O35491 1.14
soluble carrier family 2 (facilitated glucose transporter), member 9 NM_020041 1.14
muscle specific gene AF085358 1.14
cyclin T2 NM_001241 1.14
Consensus includes gb:A0245523 /FEA=EST /DB_XREF=gi:3840920 /DB_XREF=est:q25c04.x1 /CLONE=IMAGE:1869990 /UG=Hs.155572 Human clone 23801 mRNA sequi A0245523 1.14
thiosulfate sulfurtransferase (mofanase) D87292 1.14
uridine monophosphate synthetase (orotate phosphoribosyl transferase and orotidine-5'-decarboxylase) D86227 1.14

Consensus includes gb:BF877404 /FEA=EST /DB_XREF=gi:11951299 /DB_XREF=est:602087150F1 /CLONE=IMAGE-4251205 /UG=Hs.296356 Homo sapiens mRNA, cDNA	BF877404	1,14
kinase insert domain receptor (a type III receptor tyrosine kinase)	NM_002253	1,14
phosphatidylinositol-4-phosphate 5-kinase, type I, beta	U76551	1,14
CD83 antigen (activated B lymphocytes, immunoglobulin superfamily)	NM_004233	1,14
TC037 homolog	AF123539	1,14
cytochrome c oxidase subunit VIIa polypeptide 2 (liver)	NM_001865	1,14
hypothetical protein FLJ10402	NM_018085	1,14
dysbreverin, alpha	U46746	1,14
cytosolic ovarian carcinoma antigen 1	S72904	1,14
estrogen receptor 2 (ER beta)	A8006590	1,14
E2F transcription factor 4, p107/p130-binding	S75174	1,14
Consensus includes gb:U79276.1 /DEF=Human clone 23960 mRNA sequence, /FEA=mRNA /DB_XREF=gi:1710244 /UG=Hs.307238 Human clone 23960 mRNA sequence	U79276	1,14
homologous to yeast nitrogen permease (candidate tumor suppressor)	NM_006545	1,14
small nuclear ribonucleoprotein polypeptide A'	NM_003090	1,14
pyruvate dehydrogenase kinase, isoenzyme 1	NM_002610	1,14
SOM DNA binding protein	X63071	1,14
immunoglobulin lambda locus	AF047245	1,14
myeloid differentiation primary response gene (88)	U70451	1,14
hyposphosphatase 1	AF077198	1,14
tissue specific transplantation antigen P35B	U58766	1,14
POU domain, class 3, transcription factor 2	NM_005604	1,14
H2B histone family, member J	NM_003524	1,14
amyloid beta precursor protein binding protein 1, S9K0	NM_003905	1,14
hypothetical protein FLJ13149	NM_021826	1,14
spectrin, beta, non-erythrocytic 1	N92501	1,14
ataxia telangiectasia mutated (includes complementation groups A, C and D)	U82828	1,14
G protein-coupled receptor 51	AF089755	1,14
myosin, heavy polypeptide 8, cardiac muscle, alpha (cardiomyopathy, hypertrophic 1)	D00943	1,14
tumor rejection antigen (gp96) 1	NM_003289	1,14
zinc finger protein 137 (clone pH2-30)	NM_003438	1,14
SEC24 related gene family, member A (S. cerevisiae)	BE643231	1,14
VAMP (vesicle-associated membrane protein)-associated protein A (33KD)	AK000918	1,14
hypothetical protein FLJ20604	NM_017897	1,14
WD40 protein Clao1	NM_004804	1,14
SCAN domain-containing 2	AK022844	1,14
KIAA0395 protein	NM_015082	1,14
TEA domain family member 4	U63524	1,14
hypothetical protein MG35347	NM_024063	1,14
IQ motif containing GTPase activating protein 1	AI678073	1,14
calyph-phosphate memosyltransferase polypeptide 1, catalytic subunit	NM_003859	1,14
HOCMA16P protein	AA160161	1,14
HIV-1 rev binding protein 2	NM_007043	1,14
triggering receptor expressed on myeloid cells 1	NM_018643	1,14
apoptotic protease activating factor	AF149794	1,14
Consensus includes gb:AL122122.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L098 (from clone DKFZp434L098), /FEA=mRNA /DB_XREF=gi:6102949 /UG=Hs.274559 H	AL122122	1,14
inositol polyphosphate-3-phosphatase, 40KD	NM_005539	1,14
hypothetical protein	AA046752	1,14
BRCA1 associated protein	AF035620	1,14
LIM domain-containing preferred translocation partner in lipoma	BF221862	1,14
microtubule-associated protein tau	AI870749	1,14
small nuclear ribonucleoprotein polypeptide B"	NM_003092	1,14
mitochondrial ribosomal protein S14	BC254653	1,14
gb:AF132204.1 /DEF=Homo sapiens PRO2259 mRNA, complete cds, /FEA=mRNA /PROD=PRO2259 /DB_XREF=gi:11493553 /UG=Hs.302159 Homo sapiens PRO2259 mRN	AF132204	1,14
DKFZP727M1111 protein	NM_024584	1,14
hypothetical protein FLJ13648	AI268381	1,14
regulatory solute carrier protein, family 1, member 1	BF346592	1,14
KIAA0984 protein	AK000864	1,14
Consensus includes gb:AK000864.1 /DEF=Homo sapiens cDNA FLJ10002 fts, clone HEMBA1000046, /FEA=mRNA /DB_XREF=gi:7021188 /UG=Hs.296522 Homo sapiens cC	AK000864	1,14
alpha-2-glycoprotein 1, zinc	AC004522	1,14
GTF2I repeat domain-containing 1	NM_016328	1,14
hypothetical protein FLJ23514	NM_021627	1,14
CGI-43 protein	AL550875	1,14
glioma tumor suppressor candidate region gene 2	NM_015710	1,14

like mouse brain protein E48
 Consensus includes gb:BE259050 /FEA=EST /DB_XREF=gi:129548 /DB_XREF=est:8011005SF1 /CLONE=IMAGE:3350985 /UG=Hs.307137 Human DNA sequence from cl BE259050
 inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase beta
 hypothetical protein FLJ20069
 ELK1, member of ETS oncogene family
 gb:NM_025156.1 /DEF=Homo sapiens hypothetical protein FLJ12474 (FLJ12474), mRNA, /FEA=mRNA /GEN=FLJ12474 /PROD=hypothetical protein FLJ12474 /DB_XREF=gi:1025156
 T cell receptor beta locus
 SP110 nuclear body protein
 hypothetical protein PRO2389
 polyoma 1
 carboxypeptidase D
 hypothetical protein FLJ22672
 KIAA0150 protein
 GSS953 protein
 DEADH (Asp-Glu-Ala-AspHis) box binding protein 1
 integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDP2, MSK12)
 synovial sarcoma, X breakpoint 3
 bryophyllin, subfamily 2, member A2
 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsin, antitrypsin), member 7
 similar to rat myomegalin
 hypothetical protein FLJ13556
 cleavage stimulation factor, 3' pre-RNA, subunit 2, 64KD
 nitfurin 2
 Consensus includes gb:AW269792 /FEA=EST /DB_XREF=gi:6556822 /DB_XREF=est:xv4503.x1 /CLONE=IMAGE:2816069 /UG=Hs.5010 Homo sapiens clone 24672 mRNA AW269792
 poly(A)-specific ribonuclease (deadenylation nuclease)
 KIAA0556 protein
 Consensus includes gb:BG491393 /FEA=EST /DB_XREF=gi:13452905 /DB_XREF=est:602535682F1 /CLONE=IMAGE:4684748 /UG=Hs.183110 EST's, Weakly similar to ALU BG491393
 integrin beta 3 binding protein (beta3-endonexin)
 TNF receptor-associated factor 3
 hypothetical gene MGC19595
 Consensus includes gb:AK022065.1 /DEF=Homo sapiens cDNA FLJ12003 fls, clone HEMBB1001537 /FEA=mRNA /DB_XREF=gi:10433385 /UG=Hs.295683 Homo sapiens c AK022065
 centaurin, gamma 1
 HSPC016 protein
 hypothetical protein FLJ13134
 DiGeorge syndrome gene D
 ubiquitin-conjugating enzyme E2D 2 (UBCA5 homolog, yeast)
 RcoQ protein-like 5
 KIAA0728 protein
 low density lipoprotein receptor-related protein 5
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 2
 SAM domain and HD domain, 1
 Consensus includes gb:AK024543.1 /DEF=Homo sapiens cDNA: FLJ20890 fls, clone ADKA03323 /FEA=mRNA /DB_XREF=gi:10436848 /UG=Hs.306586 Homo sapiens cDNA: AK024543
 soluble carrier family 25 (mitochondrial carrier, brain), member 14
 mitogen-activated protein kinase kinase kinase 5
 gb:L14456.1 /DEF=Human Ig rearranged mu-chain gene V-N-D-N-J-region, complete cds, /FEA=mRNA /DB_XREF=gi:398497 /FL=gb:L14456.1
 hypothetical protein FLJ11787
 ATPase, Ca++ transporting, cardiac muscle, slow twitch 2
 endothelin 2
 polymerase (DNA directed), alpha
 mitogen-activated protein kinase-activated protein kinase 2
 plectin 1, intermediate filament binding protein, 500KD
 retinoblastoma binding protein 6
 KIAA1041 protein
 hemoglobin, epsilon 1
 transmembrane protein 4
 hypothetical protein MGC2470
 Consensus includes gb:AU147166 /FEA=EST /DB_XREF=gi:1008687 /DB_XREF=est:AU147166 /CLONE=HEMBB1002582 /UG=Hs.301683 Homo sapiens cDNA FLJ12091
 BRP1 homolog, subunit of RNA polymerase II (transcription initiation factor IIIB (S.cerevisiae))
 guanine nucleotide binding protein (G protein), q polypeptide
 transketolase-like 1
 SluH-interacting protein
 gamma-aminobutyric acid (GABA) A receptor, alpha 6
 misato
 NM_018116

MCM4 minichromosome maintenance deficient 4 (S. cerevisiae)	AI859865	1,14
sn3-associated polypeptide, 30kD	NM_003864	1,14
astroctadin 2	AK024064	1,14
potassium voltage-gated channel, Shaw-related subfamily, member 1	NM_004876	1,14
hypothetical protein MGC5149	U79260	1,14
KIAA0882 protein	BE646554	1,14
polymerase (RNA) II (DNA directed) polypeptide L (7.6kD)	AA772747	1,14
hypothetical protein FLJ10874	NM_018252	1,14
tailed-like kinase 2	AU151689	1,14
KIAA0494 gene product	NM_014774	1,14
hypothetical protein	BC001830	1,14
KIAA0121 gene product	D50911	1,14
myosin, light polypeptide 4, alkali; atrial, embryonic	AI133353	1,14
hepatocyte nuclear factor 4, alpha	NM_000457	1,14
DKFZP96G011 protein	AK023204	1,14
complement component 4 binding protein, alpha	NM_000715	1,14
hypothetical protein	BC001393	1,14
RNA binding protein S1, serine-rich domain	BC001659	1,14
reticulon 2	NM_005619	1,14
DnaJ (Hsp40) homolog, subfamily C, member 3	NM_006260	1,14
hypothetical protein	NM_016625	1,14
RPB5-mediated protein	AW003989	1,14
ribosomal protein L28 pseudogene 1	NM_018093	1,14
UDP-glucose ceramide glucosyltransferase	NM_003358	1,14
GS2 gene	AI814551	1,13
tripartite motif-containing 3	AA19307	1,13
NADH dehydrogenase (ubiquinone) 1, alpha/beta subcomplex, 1 (8kD, SDAP)	NM_005003	1,13
butyrophilin, subfamily 2, member A2	W58757	1,13
NADPH oxidase 3	NM_015718	1,13
HIV-1 Rev binding protein	AI742628	1,13
phospholamban	NM_002687	1,13
Consensus includes gb:AK027161.1 /DEF=Homo sapiens cDNA: FLJ23508 fs, clone LNG03158, /FEA=mRNA /DB_XREF=gi:10440220 /UG=Hs.288772 Homo sapiens cDNA: AK027161	AF18616	1,13
synaptotagmin 2	AB000359	1,13
phosphatidylinositol glycan, class C, pseudogene 1	BE966197	1,13
hypothetical protein FLJ14827	NM_018096	1,13
hypothetical protein similar to beta-transducin family	AW872377	1,13
supervillin	NM_005035	1,13
polymerase (RNA) mitochondrial (DNA directed)	AI768122	1,13
eukaryotic translation initiation factor 4 gamma, 3	NM_002770	1,13
protease, serine, 2 (trypsin 2)	NM_000559	1,13
hemoglobin, gamma A	NM_004861	1,13
cerebroside (3'-phosphoadenylylsulfate:galactosylceramide 3') sulfotransferase	BC003574	1,13
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8)	NM_002101	1,13
glycophorin C (Gerbach blood group)	NM_003563	1,13
specie-type POZ protein	AI745225	1,13
heterogeneous nuclear ribonucleoprotein R	NM_023924	1,13
KIAA0414 protein	D83077	1,13
hypothetical protein FLJ13441	AF240468	1,13
tetratropoepids repeat domain 3	AF080586	1,13
nicastin	AV713720	1,13
galenin receptor 2	BC003086	1,13
lymphocyte antigen 117	NM_018105	1,13
hypothetical protein FLJ20011	AF124491	1,13
hypothetical protein FLJ10477	AI653960	1,13
G protein-coupled receptor kinase-interactor 2	NM_015899	1,13
Consensus includes gb:AI653960 /FEA=EST /DB_XREF=gi:4737939 /DB_XREF=est:ty03h10.x1 /CLONE=IMAGE:2278051 /UG=Hs.196555 ESTs	NM_008070	1,13
putative glycolipid transfer protein	L08429	1,13
TRK-fused gene	AL572341	1,13
Rhesus blood group, D antigen	AL532341	1,13
capping protein (actin filament) muscle Z-line, beta	AA593983	1,13
ARP1 actin-related protein 1 homolog A, contractin alpha (yeast)	AI744229	1,13
SWI/SNF related, maint. associatd, actin dependent regulator of chromatin, subfamily c, member 1		
zinc finger protein 144 (Me1-16)		

KIAA1030 protein
platelet-derived growth factor receptor-like
glioma tumor suppressor candidate region gene 1
potassium voltage-gated channel, subfamily G, member 2
ATP synthase, H⁺ transporting, mitochondrial F0 complex, subunit F6
growth associated protein 43
melan-A
apoptosis regulator
hypothetical protein FLJ13490
heterogeneous nuclear ribonucleoprotein H3 (2H9)
nuclear receptor coactivator 3
COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 3
Cluster Inc. AIB32248:14910.1 Homo sapiens cDNA, 3' end /clone=IMAGE-2075692 /clone_end=3 /gp=AIB32249 /ug=Hs.211522 /len=545
KIAA0544 protein
cysteine-rich, angiogenic inducer, 61
CD59 antigen p16-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EJ32 and G344)
hypothetical protein FLJ22195
cystatin 8 (cystatin-related epididymal specific)
hypothetical protein DKFZ664O043
dUTP pyrophosphatase
hypothetical protein FLJ14124
hemopexin
cyclin G associated kinase
down-regulator of transcription 1, TBP-binding (negative cofactor 2)
IRB (bacterial acetoacetate synthase)-like
RAS27A, member RAS oncogene family
interleukin 19
spillover protein A-1
KIAA1277 protein
thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)
cysteine-rich protein 2
A kinase (PRKA) anchor protein 1
PTD18 protein
ADP-ribosylation factor 3
retinoblastoma binding protein 7
KIAA1282 protein
v-erbA sarcoma virus CT10 oncogene homolog (avian)
KIAA0853 protein
Tax interaction protein 1
amphiphysin (Shiga-Mann syndrome with breast cancer 128kD autoantigen)
associated molecule with the SH3 domain of STAM
GLUT4 enhancer factor
v-erbA sarcoma virus E28 oncogene homolog 1 (avian)
solute carrier family 21 (organic anion transporter), member 3
hypothetical protein FLJ23233
KIAA0853 protein
sorting nexin 3
T-box 21
heterogeneous nuclear ribonucleoprotein O (AU-rich element RNA binding protein 1, 37kD)
phosphoryl pyrophosphate synthetase-associated protein 1
growth hormone inducible transmembrane protein
keratin, hair, acidic38
hypothetical protein FLJ11106
potassium voltage-gated channel, KQT-like subfamily, member 2
calcium/calmodulin-dependent protein kinase (CaM kinase) II beta
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2
Consensus includes gb-BG390664 /FEA=EST /DB_XREF=gi:13284112 /DB_XREF=est:602415258F1 /CLONE=IMAGE-4523487 /UG=Hs.97496 YY1 transcription factor /FL=gb-BG390664
neutrophin 3
KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2
Rho-associated, coiled-coil containing protein kinase 1
cystatin S
mannan-binding lectin serine protease 2
BCL2-antagonist of cell death

AB028953 1.13
NM_006207 1.13
NM_015711 1.13
NM_012283 1.13
NM_001685 1.13
AF279774 1.13
NM_005511 1.13
NM_018561 1.13
NM_024942 1.13
AF132363 1.13
A1438999 1.13
NM_003653 1.13
AIB32249 1.13
AB011116 1.13
AF003114 1.13
AL513759 1.13
AL523965 1.13
NM_005492 1.13
AK001389 1.13
AB049113 1.13
NM_024868 1.13
BC005395 1.13
D88435 1.13
NM_001938 1.13
AC004794 1.13
AF125393 1.13
NM_013371 1.13
X02162 1.13
AA127623 1.13
X55005 1.13
U36190 1.13
U34074 1.13
NM_016125 1.13
NM_001659 1.13
NM_002893 1.13
AB033088 1.13
NM_016823 1.13
AB023170 1.13
AK001327 1.13
NM_001635 1.13
NM_006463 1.13
NM_020062 1.13
NM_005238 1.13
AF085224 1.13
AA429615 1.13
AL136745 1.13
BE962615 1.13
NM_013351 1.13
W74620 1.13
NM_002766 1.13
AL136713 1.13
NM_002279 1.13
NM_018324 1.13
NM_004516 1.13
U23460 1.13
NM_003070 1.13
BG390664 1.13
NM_002527 1.13
AL542253 1.13
N22548 1.13
NM_001899 1.13
AB008047 1.13
U66879 1.13

Consensus includes gb:A1694303 /FEA=EST /DB_XREF=gi:4971643 /DB_XREF=est:wd45d03.x1 /CLONE=IMAGE:2331076 /UG=Hs.15740 Homo sapiens mRNA; cDNA DKF; A1694303
steroid antigen 1
hypothetical protein FLJ23045
N-acetylglucosaminidase, alpha- (Sanfilippo disease IIIB)
RAS-related on chromosome 22
nuclear factor of kappa light polypeptide gene enhancer in B-cells 1 (p105)
engulfment and cell motility 2 (ced-12 homolog, C. elegans)
NMyc downstream-regulated gene 3
protein geranylgeranyltransferase type 1, beta subunit
Bacillus subtilis REF-X04603 /DEF=B subtilis ttrC, ttrB genes corresponding to nucleotides 995-1562 of X04603 /LEN=2073 (-5, -M, -3 represent transcriptional regions 5 prime, 3 prime, -10)
gb:BC002878.1 /DEF=Homo sapiens. Similar to collector required for Sp1 transcriptional activation, subunit 8 (34kD), clone MGC:11274, mRNA, complete cds. /FEA=mRNA /P BC002878
hypothetical protein FLJ22329
SH2 domain-containing 3A
serine racemase
nicotinamide N-methyltransferase
aldo-keto reductase family 7, member A2 (allatoin aldehyde reductase)
inactivated polyphosphate-4-phosphatase, type 1, 107kD
hypothetical protein FLJ21940
early growth response 3
hypothetical protein FLJ11370
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting, 4 (parvulin)
hypothetical protein MGC4175
adaptor-related protein complex 2, beta 1 subunit
TBC1 (tre-2/USP6, BUB2, cdc16) domain family, member 1
Consensus includes gb:AK022632.1 /DEF=Homo sapiens cDNA FLJ12570 fs, clone NT2RM4000895, /FEA=mRNA /DB_XREF=gi:10434135 /UG=Hs.142076 Homo sapiens c
zinc finger protein 174
thrombomodulin
lambda-crystallin
membrane collector protein (CD46, trophoblast-lymphocyte cross-reactive antigen)
Consensus includes gb:AL050136.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586L141 (from clone DKFZp586L141). /FEA=mRNA /DB_XREF=gi:4884346 /UG=Hs.140945 H
DKFZP564C152 protein
hypothetical protein FLJ20406
hypothetical protein FLJ13910
amyloid beta (A4) precursor protein-binding, family B, member 2 (Fe65-like)
low density lipoprotein-related protein-associated protein 1 (alpha-2-macroglobulin receptor-associated protein 1)
pancreatic polypeptide receptor 1
hypothetical protein FLJ10511
zinc finger protein 36, C3H type-like 2
glycogen synthase 1 (muscle)
G protein-coupled receptor 15
fibulin 1
hypothetical protein FLJ21562
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2
POU domain, class 5, transcription factor 1
mitochondrial/ribosomal protein S14
arrest-associated endoplasmic reticulum protein 1; ribosome associated membrane protein 4
acyl-Coenzyme A oxidase 3, pristanoyl
hypothetical protein LOC63929
hypothetical protein FLJ22709
Cluster Ind. N92708.zb44cd4.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:306438 /clone_end=3 /gb=N92708 /gi=1265017 /ug=Hs.32374 /len=668
hypothetical protein IMAGE:4215339
modulator recognition factor 1
KIAA1547 protein
protease, serine, 1 (trypsin 1)
KIAA0712 gene product
calreticulin
integrin, alpha 8
ATPase, H+ transporting, lysosomal (vacuolar proton pump), subunit 1
polymerase (RNA) II (DNA directed) polypeptide F
angiotensin I converting enzyme (peptidyl-dipeptidase A) 1
interleukin 22 receptor
parathyroid hormone-like hormone

KIAA0268 protein
GA binding protein transcription factor, alpha subunit (80kD)
calcium channel, voltage-dependent, alpha 2/delta subunit 1
zinc finger protein 302
neuroblastoma, suppression of tumorigenicity 1
general transcription factor IIIC, polypeptide 1 (alpha subunit, 220kD)
serine/threonine kinase 17a (apoptosis-inducing)
serine palmitoyltransferase, long chain base subunit 2
Hiv-1 inducer of short transcripts binding protein; lymphoma related factor
HEMk homolog 7/b
arginine vasopressin receptor 2 (nephrogenic diabetes insipidus)
KIAA0494 gene product
hypothetical gene supported by AF038182; BC009203
growth arrest-specific 6
protection of telomeres 1
transposon-derived Butler1 transposase-like protein
FH/IFH2 domain-containing protein
RAN binding protein 17
CGI-74 protein
BRF2, subunit of RNA polymerase III transcription initiation factor, BRF1-like
hypothetical protein MG22654
Consensus includes gb:AL050002.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564O222 (from clone DKFZp564O222). /FEA=mRNA /DB_XREF=gi:4884256 /UG=Hs.94795
ARP2 actin-related protein 2 homolog (yeast)
oxidative membrane protein
decorin
OCS3 general control of amino-acid synthesis 5-like 2 (yeast)
Consensus includes gb:A1923972 /FEA=EST /DB_XREF=gi:5659936 /DB_XREF=est:wn49c10.x1 /CLONE=IMAGE:2448786 /UG=Hs.127723 Homo sapiens mRNA; cDNA DKFZp564O222 (from clone DKFZp564O222). /FEA=mRNA /DB_XREF=gi:4884256 /UG=Hs.94795
hypothetical protein R30953_1
chromosome 17 open reading frame 1A
Wiskott-Aldrich syndrome protein interacting protein
Consensus includes gb:AK000270.1 /DEF=Homo sapiens cDNA FLJ20263 fs, clone COLF7804, highly similar to AJ131693 Homo sapiens mRNA for AKAP450 protein. /FEA=AK000270
interferon, gamma
KIAA1069 protein
mesothionein 1X
KIAA0915 gene product
hypothetical protein FLJ10468
plaiophilin 4
ubiquitin-activating enzyme E1-like
hypothetical protein
KIAA0409 protein
Meis1, myeloid ecotropic viral integration site 1 homolog 2 (mouse)
calpain 7
RAD21 homolog (S. pombe)
hypothetical protein
CCAA1/henancer binding protein (C/EBP), delta
Consensus includes gb:Z98950 /DEF=Human DNA sequence from PAC 507115 on chromosome Xq28.3-27.3. Contains 605 ribosomal protein L44 (L41, L36) like gene. ESTs, A1982377
PTPRF interacting protein, binding protein 1 (liprin beta 1)
cadherin 16, XSP-cadherin
Kruppel-like factor 4 (gut)
calcium channel, voltage-dependent, alpha 1 subunit
hypothetical protein FLJ12496
integrin, alpha 4 (enigen CD49D, alpha 4 subunit of VLA-4 receptor)
meth homo box homolog 2 (Drosophila)
laminin, beta 3 (lncin (125kD), kalinin (140kD), BM600 (125kD))
phosphatidylinositol-3 phosphatase adaptor subunit
hypothetical protein FLJ22222
casein kinase 1, alpha 1
interleukin 1, alpha
CD58 antigen, (lymphocyte function-associated antigen 3)
HNDEL-iso protein
U6 snRNA-associated Sm-like protein
synapobin 2
Consensus includes gb:AK02679.1 /DEF=Homo sapiens cDNA FLJ12617 fs, clone NT2RM4001662. /FEA=mRNA /DB_XREF=gi:10434213 /UG=Hs.287538 Homo sapiens c AK02679

CD28 antigen (Tp44)
 ferritin, heavy polypeptide 1
 myosin 5C
 HUO1 gene
 ribosomal protein S29
 glycosylphosphatidylinositol specific phospholipase D1
 Consensus includes gp:AL049037 /FEA=EST /DB_XREF=gi:4728346 /DB_XREF=est:DKFZp434P1518_r1 /CLONE=DKFZp434P1518 /UG=Hs.323833 syntaphilin
 PR domain containing 4
 hypothetical C2H2 zinc finger protein FLJ22504
 hypothetical protein FLJ20150
 Consensus includes gp:AK021982.1 /DEF=Homo sapiens cDNA FLJ11930 fls, clone HEMBB100044.1 /FEA=mRNA /DB_XREF=gi:10433301 /UG=Hs.308618 Homo sapiens c AK021992
 beaded filament structural protein 2, plakophilin
 hypothetical protein MQC2821
 Consensus includes gp:AL14252 /FEA=EST /DB_XREF=gi:5425467 /DB_XREF=est:w70g04.x1 /CLONE=IMAGE:2408214 /UG=Hs.194051 Homo sapiens mRNA; cDNA DKF A1814252
 leukotriene b4 receptor (chemokine receptor-like 1)
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 7
 perlecanin material 1
 P311 protein
 hypothetical protein FLJ12438
 RAN binding protein 2
 synovial sarcoma translocation, chromosome 18
 15 kDa selenoprotein
 tyrosine kinase, non-receptor, 1
 core promoter element binding protein
 hypothetical protein FLJ11132
 RNA binding motif protein 15
 chondroitin sulfate proteoglycan 4 (melanoma-associated)
 TRAM-like protein
 hypothetical protein DKFZp434E2135
 CD2 antigen (cytoplasmic tail) binding protein 2
 hypothetical protein MQC2508
 protein tyrosine phosphatase type IVA, member 3
 retinoblastoma (X-linked, juvenile) 1
 methylenetetrahydrofolate dehydrogenase
 mitogen-activated protein kinase 3
 apolipoprotein E
 TEA domain family member 1 (SV40 transcriptional enhancer factor)
 farnesyltransferase, CAAAX box, alpha
 ubiquitin-conjugating enzyme E2C
 spermatogenesis associated 2
 hypothetical protein MQC4126
 soluble carrier family 19 (folate transporter), member 1
 ribosomal protein S20
 Consensus includes gp:D16471.1 /DEF=Human mRNA, Xq terminal portion. /FEA=mRNA /DB_XREF=gi:495089 /UG=Hs.121571 Human mRNA, Xq terminal portion
 vascular endothelial growth factor
 pyridine-5-carboxylate reductase 1
 homeo box 11-like 2
 hypothetical protein MQC4178
 cylindromatosis (turban tumor syndrome)
 40S ribosomal protein S27 isoform
 growth differentiation factor 11
 wingless-type MMTV integrallon site family, member 6
 opoid receptor, kappa 1
 endocytic receptor (macrophage mannose receptor family)
 SH3-domain binding protein 2
 glioma amplified on chromosome 1 protein (leucine-rich)
 KIAA0733 protein
 cyclic nucleotide gated channel beta 3
 vitiligo-associated protein VIT-1
 phospholipase C, epsilon 2
 transmembrane 4 superfamily member 8
 Consensus includes gp:AL023584 /DEF=Human DNA sequence from clone 67K17 on chromosome 6q24.1-24.3. Contains the HIVEP2 (Schnurrl-2) gene for HIV type 1 Enhanc
 M10098 Human 18S rRNA sequence, length 1869 bases, 5 prime target bases 1-646

hypothetical protein DKFZ0434.037
 Consensus includes gb:AA148507 /FEA=EST /DB_XREF=gi:1721551 /DB_XREF=est:z06005.s1 /CLONE=IMAGE:491528 /UG=Hs.14846 Homo sapiens mRNA; cDNA DKFZ: AA148507
 zinc finger protein 157 (ZFZ22)
 NM_003446 1.11
 insulin (Strongylocentrotus purpuratus) homolog 2 (actin-bundling protein, retinal)
 NM_012418 1.11
 AK001574
 NM_007260 1.11
 gng1 phosphoprotein 5
 NM_014347 1.11
 lysophospholipase II
 NM_006346 1.11
 progesterone-induced blocking factor 1
 NM_005355 1.11
 kinesin-like 3
 AF268617 1.11
 POU domain, class 5, transcription factor 1
 AU168371 1.11
 homeo box D1
 Consensus includes gb:AU145746 /FEA=EST /DB_XREF=gi:1007267 /DB_XREF=est:AU145746 /CLONE=HEMBA1005679 /UG=Hs.82193 esterase Olfmlyglutathione hydrolase
 NM_004231 1.11
 ATPase, vesicular, 14 kD
 NM_012414 1.11
 rab3 GTPase-activating protein, non-catalytic subunit (150kD)
 NM_000515 1.11
 growth hormone 1
 BF223718 1.11
 titin functional domain (PTPRF interacting)
 NM_006839 1.11
 inner membrane protein, mitochondrial (mitofilin)
 NM_003039 1.11
 solute carrier family 2 (facilitated glucose/fructose transporter), member 5
 NM_030758 1.11
 cysteine binding protein 2
 Consensus includes gb:AL078598 DEF=Human DNA sequence from clone RP3-42905 on chromosome 6q21-22.1. Contains the NR2E1 gene for nuclear receptor 2E1 (allies AL078598)
 NM_002599 1.11
 phosphodiesterase 2A, cGMP-stimulated
 AF127765 1.11
 calpain 3, (p94)
 BF213829 1.11
 D78261 1.11
 interferon regulatory factor 4
 NM_002041 1.11
 GA binding protein transcription factor, beta subunit 2 (47kD)
 NM_006810 1.11
 for protein disulfide isomerase-related
 AB010153 1.11
 tumor protein 63 kDa with strong homology to p53
 NM_024673 1.11
 hypothetical protein FLJ12270
 NM_061065 1.11
 tumor necrosis factor receptor superfamily, member 1A
 M87789 1.11
 immunoglobulin heavy constant gamma 3 (G3m marker)
 NM_010153 1.11
 guanine nucleotide binding protein (G protein), gamma 7
 BF432795 1.11
 tumor necrosis factor, alpha-induced protein 3
 A1736896 1.11
 Consensus includes gb:AK024135.1 /DEF=Homo sapiens cDNA FLJ14073 fts, clone HEMBB1001812 /FEA=mRNA /DB_XREF=gi:10436441 /UG=Hs.296751 Homo sapiens cAK024135
 v-ort avian sarcoma virus CT10 oncogene homolog
 NM_003931 1.11
 WAS protein family, member 1
 NM_019853 1.11
 protein phosphatase 4 regulatory subunit 2
 A4336566 1.11
 thionotin interacting protein
 AB25734 1.11
 Consensus includes gb:AB25734 /FEA=EST /DB_XREF=gi:5661698 /DB_XREF=est:wo34q08.x1 /CLONE=IMAGE:2457278 /UG=Hs.110613 KIAA00220 protein
 polymerase (RNA) III (DNA directed) (39kD)
 NM_006466 1.11
 calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma
 AU933569 1.11
 KIAA1018 protein
 NM_014967 1.11
 hypothetical protein MGC5363
 NM_024064 1.11
 CD22 antigen
 NM_001771 1.11
 F-box only protein 2
 NM_012168 1.11
 homeo box C11
 NM_014212 1.11
 HSPC055 protein
 NM_041453 1.11
 hypothetical protein PP1044
 NM_021730 1.11
 baculoviral IAP repeat-containing 3
 U37546 1.11
 neutral sphingomyelinase (N-SMase) activation associated factor
 NM_003580 1.11
 phosphodiesterase 4D interacting protein (myomegalin)
 AB042555 1.11
 small inducible cytokine A1, I-309
 NM_002981 1.11
 trinucleotide repeat containing 4
 NM_007185 1.11
 crystallin, zeta (quinone reductase)-like 1
 NM_005111 1.11
 hypothetical protein FLJ10700
 NM_018182 1.11
 KIAA0009 gene product
 BF214329 1.11
 telomodin 1 (smooth muscle)
 BC001755 1.11
 activating transcription factor 6
 NM_007346 1.11
 mitochondrial ribosomal protein L13
 NM_014078 1.11
 ubiquitin-conjugating enzyme E2A (RAD6 homolog)
 NM_003336 1.11
 interferon, gamma-inducible protein 16
 BG256877 1.11
 mitochondrial ribosomal protein S22
 NM_020191 1.11
 CGI-130 protein
 BC001671 1.11
 Consensus includes gb:AW274747 /FEA=EST /DB_XREF=gi:5661777 /DB_XREF=est:tn34e08.x1 /CLONE=IMAGE:2665622 /UG=Hs.270527 Homo sapiens mRNA; cDNA Df: AW274747

hypothetical protein ASH1	NM_018469	1.11
Consensus includes gb:AU144284 /FEA=EST /DB_XREF=gi:1005805 /DB_XREF=est:AU144284 /CLONE=HEMBA1001439 /UG=Hs.11801 Interferon regulatory factor 6 /FL= AU144284	1.11	
transformation/transcription domain-associated protein	NM_003496	1.11
brain link protein-1	NM_021817	1.11
Consensus includes gb:AK021960.1 /DEF=Homo sapiens cDNA FLJ11898 fs. clone HEMBA1007322. /FEA=mRNA /DB_XREF=gi:10433269 /UG=Hs.313509 Homo sapiens c AK021960	1.11	
translocase of outer mitochondrial membrane 70 homolog A (yeast)	BC003633	1.11
KIAA1089 protein	AB029012	1.11
CDC10 cell division cycle 10 homolog (S. cerevisiae)	AU157515	1.11
hypothetical protein	BC000979	1.11
Consensus includes gb:BE549732 /FEA=EST /DB_XREF=gi:9791424 /DB_XREF=est:7638a04 x1 /CLONE=IMAGE:3230478 /UG=Hs.112158 Homo sapiens PAC clone RP4-7 BE549732	1.11	
aphidopine kinase type 2 isoform	AF245447	1.11
gb:BC002361.1 /DEF=Homo sapiens. Similar to transducin (beta)-like 3. clone MGC:8613. mRNA, complete cds. /FEA=mRNA /PROD=Similar to transducin (beta)-like 3 /DB_XREF=gi:10433269 /UG=Hs.11801 Interferon regulatory factor 6 /FL= AU144284	1.11	
SHP2 interacting transmembrane adaptor	NM_014450	1.11
centaurin, delta 1	AK023737	1.11
KIAA1033 protein	AK001728	1.11
serine palmitoyltransferase, long chain base subunit 1	AL568804	1.11
gelsolin (amyloidosis, Finnish type)	NM_000177	1.11
hypothetical protein FLJ10233	NM_018034	1.11
major histocompatibility complex, class II, DP beta 1	NM_002121	1.11
DKFZPS47E1010 protein	NM_015607	1.11
RNA binding motif, single stranded interacting protein 1	D28482	1.11
DKFZP434H132 protein	NM_015492	1.11
Consensus includes gb:AW873544 /FEA=EST /DB_XREF=gi:8007597 /DB_XREF=est:th62a01.x1 /CLONE=IMAGE:3041928 /UG=Hs.277500 Human mRNA for T cell receptor AW873544	1.11	
hypothetical protein FLJ11099	NM_018320	1.11
metallothionein 1L	NM_002450	1.11
prostaglandin E receptor 4 (subtype EP4)	AI675173	1.11
aridine homolog 2 (Drosophila)	BE984655	1.11
transcription factor-like 4	AW074630	1.11
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) 9kD	BE984655	1.11
chromosome 18 open reading frame 1	AI34506	1.11
lactoferrin rich protein	NM_003945	1.11
nucleophosmin (nucleolar phosphoprotein B23, numatrin)	NM_007244	1.11
KIN, antigenic determinant of recA protein homolog (mouse)	BC002398	1.11
palate, lung and nasal epithelium carcinoma associated	NM_012311	1.11
phospholipase A2, group IVB (cytosolic)	NM_016583	1.11
KIAA0170 gene product	AK000550	1.11
transcription elongation factor B (SIII), polypeptide 1-like	NM_014641	1.11
trypsin-like peptidase II	BE984043	1.11
Sac domain-containing inositol phosphatase 2	AW612376	1.11
Consensus includes gb:AK000787.1 /DEF=Homo sapiens cDNA FLJ20780 fs. clone COL04256. /FEA=mRNA /DB_XREF=gi:7021089 /UG=Hs.157392 Homo sapiens cDNA FI AK000787	1.11	
foslin 2	BC003683	1.11
alpha-1-microglobulin/bikunin precursor	NM_001633	1.11
immunoglobulin kappa variable 1-13	AW408194	1.11
gene from NF2/meringoma region of 22q12	AB023200	1.11
methylmalonate-semialdehyde dehydrogenase	NM_005589	1.11
interleukin 5 receptor, alpha	NM_000584	1.11
mitogen-activated protein kinase kinase 1	AI571419	1.11
forhead box O3A	NM_001455	1.11
baculoviral IAP repeat-containing 4	U32974	1.11
NADH dehydrogenase (ubiquinone) Fc-S protein 4 (18kD) (NADH-coenzyme Q reductase)	BC005270	1.11
hypothetical protein FLJ10415	NM_018089	1.11
Consensus includes gb:AL080132.1 /DEF=Homo sapiens mRNA, cDNA DKFZP434F083 (from clone DKFZP434F083). /FEA=mRNA /DB_XREF=gi:5262572 /UG=Hs.118207 H AL080132	1.11	
UDP-galactose transporter related	NM_005827	1.11
hypothetical protein DKFZP434I1916	AL136771	1.11
trypsin, alpha	NM_003293	1.11
zinc finger protein 194 (Kruppel-like)	AB111577	1.11
cytochrome b-561	AL514271	1.11
heme oxygenase (decycling) 2	D21243	1.11
mitogen-activated protein kinase 4	BF115223	1.11
LAG1 longevity assurance homolog 2 (S. cerevisiae)	AK001105	1.11
KIAA0708 protein	AB014608	1.11
c-myc binding protein	AL525412	1.11
huntingtin (Huntington disease)	NM_002111	1.11

somatostatin receptor 3	NM_001051	1.11
dictyopl homolog 4 (Xenopus laevis)	NM_014420	1.11
ephrin-B1	NM_004429	1.11
transcription factor 8 (represses interleukin 2 expression)	AI061174	1.11
COQ7 coenzyme Q7 homolog ubiquinone (yeast)	AF032900	1.11
Bacillus subtilis RREF-L38424 /DEF-F8 subtilis dppB, joiF, joiG genes corresponding to nucleotides 2634-3089 of L38424 (LEN=1931 (-5, -M, -3) represent transcript regions 5' pL38424		1.11
Consensus includes gb:AF305057 /DEF=Homo sapiens RTS (RTS) gene, complete cds, alternatively spliced /FEA=mRNA_1 /DB_XREF=gi:17094017 /UG=Hs.160433 rTS bet AF305057		1.11
ELG protein	NM_018553	1.11
CASP8 and FADD-like apoptosis regulator	AF041461	1.11
CD2-associated protein	NM_012120	1.11
branched chain keto acid dehydrogenase E1, beta polypeptide (maple syrup urine disease)	AL531533	1.11
SET domain, bifurcated 1	NM_012432	1.11
Consensus includes gb:AJ39732 /FEA=EST /DB_XREF=gi:4076659 /DB_XREF=est:q42003.x1 /CLONE=IMAGE:1935201 /UG=Hs.309763 ESTs, Weakly similar to ALU7_HL AJ39732		1.10
chloride channel Kc	NM_004070	1.10
hypothetical protein FLJ23420	NM_025061	1.10
spastic paraplegia 4 (autosomal dominant; spastin)	AB076006	1.10
S100 calcium binding protein A1	NM_006271	1.10
transcription factor-like 5 (basic helix-loop-helix)	AL035689	1.10
DKFZP586G011 protein	AK021613	1.10
three prime repair exonuclease 1	BC002903	1.10
mitochondrial ribosomal protein S35	NM_021821	1.10
myelin basic protein	AW070431	1.10
tyrosyl-tRNA synthetase 2 (mitochondrial)	NM_015938	1.10
transmembrane 9 superfamily member 1	BE899402	1.10
ectonucleoside triphosphate diphosphohydrolase 1	AV717590	1.10
hydroxysteroid (17-beta) dehydrogenase 1	NM_000413	1.10
OtaJ (Hsp40) homolog, subfamily B, member 4	NM_007034	1.10
hypothetical protein FLJ12505	NM_024749	1.10
postrypsinic protein CRIPT	NM_014171	1.10
cyclochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 7, pseudogene 2	U22044	1.10
PCTAIRE protein kinase 1	BC001048	1.10
pannexin 1	NM_015388	1.10
semaphorin domain, immunoglobulin domain (Ig), transmembrane domain TM) and short cytoplasmic domain, (semaphorin) 4C	NM_017789	1.10
amyloid beta (A4) precursor-like protein 2	NM_001642	1.10
hypothetical protein DKFZp434O1427	NM_030963	1.10
HOG-like suppressor	NM_020386	1.10
mitogen-activated protein kinase-activated protein kinase 5	NM_003668	1.10
ring finger protein 24	NM_022823	1.10
hypothetical protein FLJ22362	BC000213	1.10
transcription factor-like 5 (basic helix-loop-helix)	NM_008602	1.10
KRAB-zinc finger protein SZF1-1	AF114817	1.10
uracil-DNA glycosylase pseudogene 2	X82986	1.10
v-yes-1 (Yamaguchi) sarcoma viral related oncogene homolog	AI356412	1.10
frequently rearranged in advanced T-cell lymphomas	NM_005479	1.10
erythrin 4A (placental)	NM_004604	1.10
Int D proto-oncogene	NM_005354	1.10
HCF-binding transcription factor Zhangfei	AW204564	1.10
nucleoporin 62KD	NM_012346	1.10
Cluster Ind. AJ302185:qr55d02.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:1902435 /clone_end=3' /gb=AI302185 /gi=3961531 /ug=Hs.4789 /len=522	AI302185	1.10
keratoprotein alpha 8 (keratin alpha 7)	AI002111	1.10
CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EL32 and G344)	X16447	1.10
HSPC932 protein	NM_014150	1.10
KIAA0283 gene product	BE614199	1.10
microtubule-interacting protein that associates with TRAF3	AF230877	1.10
SEC23 protein	AK001465	1.10
dysferlin 1, longin (autosomal dominant; torsin A)	BC000674	1.10
expression of tumorigenicity 13 (colon carcinoma) (Hsp70 interacting protein)	U17714	1.10
tubulin-like 1	AI769310	1.10
hypothetical protein FLJ10111	NM_017999	1.10
nuclear transcription factor Y, beta	AI804118	1.10
melanoma antigen, family B, 1	NM_002363	1.10
KIAA1668 protein	W46406	1.10
Consensus includes gb:AA133342 /FEA=EST /DB_XREF=gi:1690310 /DB_XREF=est:zn92n03.s1 /CLONE=IMAGE:585685 /UG=Hs.6700 ESTs	AA133342	1.10

hypothetical protein FLJ20450
 KIAA0931 protein
 HIRA interacting protein 3
 ferredoxin reductase
 hypothetical protein FLJ20644
 hypothetical protein FLJ20188
 cholinergic receptor, nicotinic, alpha polypeptide 2 (neuronal)
 KIAA1284 protein
 testis expressed sequence 13A
 nardilysin (N-angiotensin diester convertase)
 hypothetical protein FLJ10382
 gpA1027523.1 IDEF+Homo sapiens transcription factor T2P (T2P) mRNA, complete cds. /FEA=mRNA /GEN=T2P /PROD=transcription factor T2P /DB_XREF=gi:13195150 /A AY027523
 set-1 suppressor of lin-12-like (C. elegans)
 collagen, type XVIII, alpha 1
 phosphatidylinositol transfer protein
 interleukin 16 (lymphocyte chemoattractant factor)
 proliferation-associated 204, 38kD
 Consensus includes gpA1137398.1 IDEF+EST /DB_XREF=gi:4188641 /DB_XREF=est:tc87b04.x1 /CLONE=IMAGE:2069647 /UG=Hs.153445 Human mRNA for unknown product /A1378788
 lymphocyte antigen 17
 Consensus includes gpA1137398.1 IDEF+Homo sapiens mRNA, cDNA DKFZp434K052 (from clone DKFZp434K052). /FEA=mRNA /DB_XREF=gi:6807944 /UG=Hs.169639 /A137398
 sialyltransferase 8 (alpha-2, 8-polysialyltransferase) D
 solute carrier family 6 (neurotransmitter transporter, L-proline), member 7
 hypothetical protein FLJ14827
 KIAA0767 protein
 nucleotide-sugar transporter similar to C. elegans sqv-7
 GTP-binding protein ragB
 KIAA0916 protein
 jagged 1 (Alagille syndrome)
 chromosome 2 open reading frame 6
 dystrobrevin, alpha
 ref1004
 Consensus includes gpA1041204 /FEA=EST /DB_XREF=gi:3280398 /DB_XREF=est:ov77g06.x1 /CLONE=IMAGE:1643386 /UG=Hs.323748 Homo sapiens clone CDABP0086 /A041204
 solute carrier family 6 (neurotransmitter transporter, glycine), member 9
 breast carcinoma amplified sequence 4
 COI-02 protein
 ATP-binding cassette, sub-family C (CFTR/MRP), member 4
 hypothetical protein FLJ23445
 dishevelled, dish homolog 3 (Drosophila)
 xenoderm pigmentosum, complementation group C
 regulator of G-protein signalling 1
 gpNM_004692.1 IDEF+Homo sapiens internerxin neuronal intermediate filament protein, alpha (INA). mRNA. /FEA=mRNA /GEN=INA /PROD=interneuron neuronal intermediate filament protein /A04692
 putative human HLA class II associated protein 1
 KIAA0576 protein
 suppressor of Ty 8 homolog (S. cerevisiae)
 solute carrier family 24 (sodium/potassium/calcium exchanger), member 1
 hypothetical protein DKFZp564A022
 cAMP responsive element modulator
 dimethylarginine dimethylaminohydrolase 2
 TNFRSF1A-associated via death domain
 LIM domain only 6
 gpL00961.1 IDEF+Homo sapiens transmembrane tyrosine kinase mRNA, complete cds. /FEA=CDs /PROD=tyrosine kinase /DB_XREF=gi:897614 /FL=gpL00961.1
 syntaxin 3A
 neogenin homolog 1 (chicken)
 BWR1 protein
 X transporter protein 3
 mucosal vascular addressin cell adhesion molecule 1
 thyroid hormone receptor-associated protein, 95-kD subunit
 transcription factor 1, hepatic; LF-B1, hepatic nuclear factor (HNF1), albumin proximal factor
 hypothetical protein FLJ22055
 syntaxin binding protein 2
 baculoviral IAP repeat-containing 5 (sunvin)
 UBX domain-containing 1
 KIAA0463 protein

hypothetical protein FLJ14213	NM_024841	1.10
hypothetical protein FLJ23132	NM_025054	1.10
KIA0329 gene product	NM_014844	1.10
lifeguard	AB023167	1.10
hypothetical protein FLJ12671	AW294567	1.10
MYLE protein	NM_014015	1.10
tumor necrosis factor receptor superfamily, member 10b	AF016266	1.10
A kinase (PRKA) anchor protein 10	A4456929	1.10
nuclear respiratory factor 1	L22454	1.10
phosphoribosyl pyrophosphate synthetase 1	BC001605	1.10
Glutamate dehydrogenase-2	AC006144	1.10
SFRS protein kinase 2	AC005070	1.10
insulin receptor-related receptor	J05048	1.10
protein phosphatase	NM_016364	1.10
2'-5'-oligoadenylate synthetase 2 (69-71 kD)	NM_002535	1.10
NAADH oxidase 1	NM_013955	1.10
iodide receptor 1	AL050262	1.10
dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS206, DXS230, DXS239, DXS268, DXS270, DXS272	NM_004010	1.10
Reg C protein	NM_001615	1.10
peptidylglycine alpha-amidating monooxygenase COOH-terminal interactor	NM_022157	1.10
pre-mRNA splicing SR protein RA4	AF056209	1.10
Consensus includes gb:AF035278.1 /DEF+Homo sapiens clone 23676 mRNA sequence, /FEA=mRNA /DB_XREF=gi:2661027 /UG=Hs.100841 Homo sapiens clone 23676 mf	AA646521	1.10
SUMO-1 activating enzyme subunit 2	NM_005499	1.10
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 7 (RNA helicase, 52kD)	NM_004940	1.10
Consensus includes gb:AL049288.1 /DEF+Homo sapiens mRNA; cDNA DKFZp564G103 (from clone DKFZp564G103), /FEA=mRNA /DB_XREF=gi:4500017 /UG=Hs.2026841	AL049268	1.10
chromosome 11 open reading frame 13	NM_003475	1.10
sphingosine kinase type 2 isoform	AA483440	1.10
N-acetylglucosaminidase, alpha-	M38083	1.10
ADP-ribosylation factor-like 7	BC435404	1.10
Consensus includes gb:BE673445 /FEA+EST /DB_XREF=gi:10033986 /DB_XREF=est:7d35d10.x1 /CLONE=IMAGE:3249235 /UG=Hs.22049 Homo sapiens chromosome 19,	BE673445	1.10
protein kinase C-like 1	NM_002741	1.10
copine III	NM_003909	1.10
hypothetical protein FLJ12085	NM_022771	1.10
Importin alpha 6 (Importin alpha 7)	AU154321	1.10
nuclear domain 10 protein	BC004130	1.10
lectin 2 (testicular)	AB033823	1.10
p21(CDK1A)-activated kinase 4	NM_005884	1.10
dystrophia myotonia-containing WD repeat motif	L19267	1.10
hormo box B1	NM_002144	1.10
Consensus includes gb:A010895 /FEA+EST /DB_XREF=gi:3630631 /DB_XREF=est:wd20n09.x1 /CLONE=IMAGE:2328737 /UG=Hs.19720 Homo sapiens mRNA full length in	AI910895	1.10
putative methyltransferase	AW069290	1.10
KIA03596 protein	AI674759	1.10
ATP-binding cassette, sub-family C (CFTRMRP), member 1	AI539710	1.10
dystrophin related protein 2	NM_001939	1.10
hypothetical protein MG3731	AW002578	1.10
mitogen-activated protein kinase kinase kinase 1	NM_024675	1.10
SUMO-1-specific protease	NM_007161	1.10
KIA00648 protein	AF306508	1.10
mitochondrial carrier family protein	AK021757	1.10
hypothetical protein FLJ10647	NM_018843	1.10
cardioembryonic antigen-related cell adhesion molecule 8	BC006241	1.10
Down syndrome critical region gene 1-like 2	M33326	1.10
apoptosis-related protein PNAS-1	NM_013441	1.10
HUBA-inducible	NM_016534	1.10
dystrophia myotonia-protein kinase	NM_006460	1.10
Consensus includes gb:AL390738 /DEF+Human DNA sequence from clone RP11-438F9 on chromosome 13 Contains the gene for the ortholog of mouse homeobox protein G-	L08835	1.09
chorionic gonadotropin hormone 2	AL390738	1.09
low density lipoprotein receptor (familial hypercholesterolemia)	NM_022645	1.09
ubiquitin specific protease 9, X chromosome (Drosophila fat facets related)	NM_000527	1.09
Cas-Bi-M (murine) ecotropic retroviral transforming sequence	AA824386	1.09
	NM_005188	1.09

thrombosane A2 receptor
 major histocompatibility complex, class II, DM alpha
 NK2 transcription factor homolog B (Drosophila)
 Consensus includes gp:AV303478/FEA-EST/DB_XREF=gi:8713165/DB_XREF=est:zv19e10.x1/CLONE=IMAGE2813610/UG=Hs.132299 ESTs
 ghorylita reductase/hydroxypyruvate reductase
 testis expressed sequence 13B
 ATP-binding cassette, sub-family B (MDR/TAP), member 6
 collagen, type VI, alpha 1
 offactory receptor, family 2, subfamily N, member 1 pseudogene
 Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2
 hypothetical protein FLJ13119
 M11167 Human 28S rRNA sequence, length 5025 bases, middle target bases 1668-3330
 Consensus includes gbl:10374.1/DEF=Human (clone CTG-A4) mRNA sequence. /FEA=mRNA/DB_XREF=gi:307286/UG=Hs.281587 Human (clone CTG-A4) mRNA sequen
 G protein-coupled receptor, family C, group 5, member B
 endothelin type b receptor-like protein 2
 acute carrier family 4, sodium bicarbonate cotransporter, member 8
 TP53 target gene 1
 hypothetical protein FLJ10849
 cystatin F (leukocystatin)
 Consensus includes gb:A620911/FEA-EST/DB_XREF=gi:4630037/DB_XREF=est:u05412.x1/CLONE=IMAGE2250167/UG=Hs.205125 ESTs
 LIM protein (similar to rat protein kinase C-binding enigma)
 gb:L21758.1/DEF=Homo sapiens acute myeloid leukemia associated protein (AML1EAP) mRNA, complete cds. /FEA=mRNA/GEN=AML1EAP/DB_XREF=gi:400340/FL=gb:L21758
 hypothetical protein FLJ20424
 KIAA0397 gene product
 growth hormone 1
 HMB-40 nucleole
 RAB3 GTPase-ACTIVATING PROTEIN
 allograft inflammatory factor 1
 peptidylprolyl isomerase E (cyclophilin E)
 MYC-associated zinc finger protein (purine-binding transcription factor)
 N-acetyltransferase 2 (erythrine N-acetyltransferase)
 hypothetical protein PRO0871
 hypothetical protein MGC11349
 T-cell, immune regulator 1
 sodium bicarbonate transporter 4
 a disintegrin and metalloproteinase domain 28
 rat finger protein 2
 Consensus includes gb:AB050195.1/DEF=Homo sapiens mRNA, partial cds, clone H2-5, MJD homolog. /FEA=mRNA/DB_XREF=gi:11559487/UG=Hs.302079 Homo sapiens
 phosphodiesterase 8A
 potassium voltage-gated channel, Shaw-related subfamily, member 3
 hypothetical protein similar to ankyrin repeat-containing protein AKR1
 hypothetical protein FLJ20533
 hypothetical protein FLJ22643
 solute carrier family 22 (organic cation transporter), member 1-like
 fortheed box Q3B
 putative UDP-GalNAc:polypeptide N-acetylglucosaminyltransferase T9
 peptidylprolyl isomerase (cyclophilin)-like 2
 glucagon receptor
 Consensus includes gb:AC004490/DEF=Homo sapiens chromosome 19, cosmid R29381/FEA=CDS/DB_XREF=gi:2995576/UG=Hs.156015 Homo sapiens chromosome 19,
 CGI-138 protein
 SH3-domain protein 5 (ponatin)
 arbutin alien mating type information regulation 2 homolog 5 (S. cerevisiae)
 oplate receptor-like 1
 lipase, hormone-sensitive
 type I transmembrane receptor (seizure-related protein)
 95 kDa retinoblastoma protein binding protein
 BAI1-associated protein 3
 apoptosis inhibitor 5
 Consensus includes gb:AL049314.1/DEF=Homo sapiens mRNA; cDNA DKFZ564C156 (from clone DKFZ564C156). /FEA=mRNA/DB_XREF=gi:4500088/UG=Hs.2785921/AL049314
 phosphatase kinase, alpha 2 (liver)
 serine protease inhibitor, Kunitz type 1
 ubiquitin-conjugating enzyme E2D 2 (UBC4/5 homolog, yeast)
 cytochrome P450 retinoid metabolizing protein

MOST-1 protein
 hypothetical protein FLJ13330
 proteasome (prosome, macropain) subunit, alpha type 4
 hypothetical protein FLJ23429
 Consensus includes gb:AL339602.1 /DEF=Homo sapiens mRNA; cDNA DKFp547D148 (from clone DKFp547D148). /FEA=mRNA /DB_XREF=gi:8655670 /UG=Hs.306514 t AL339602
 perlecanin material 1
 arrestin 3, retinal (k-arrestin)
 histone acetyltransferase
 KIAA0553 protein
 exonuclease NEF-4p
 hypothetical protein FLJ10979
 Consensus includes gb:AK026803.1 /DEF=Homo sapiens cDNA, FLJ23150 fts, clone LNG09338. /FEA=mRNA /DB_XREF=gi:10439742 /UG=Hs.277178 Homo sapiens cDNA; BC006116
 tubulin-specific chaperone c
 hypothetical protein FLJ11722
 Consensus includes gb:AF258545 /DEF=Homo sapiens chromosome 17, HC90, HC71AC, HC8 and HC56 genes, complete sequence /FEA=CDS /DB_XREF=gi:10945429 /UG=Hs.326457 ESTs
 spindle pole body protein
 cell division cycle 25C
 SRY (sex determining region Y)-box 2
 KIAA1862 protein
 nuclear receptor subfamily 2, group F, member 8
 B-cell CLL/lymphoma 7C
 interleukin 17 receptor
 single Ig IL-1R-related molecule
 KIAA1033 protein
 Consensus includes gb:AA648851 /FEA=EST /DB_XREF=gi:2577179 /DB_XREF=est:ns53g06.s1 /CLONE=IMAGE:1187386 /UG=Hs.326457 ESTs
 PR domain containing 4
 GNAS complex locus
 PDZ-binding kinase; T-cell originated protein kinase
 hypothetical protein FLJ21945
 colony stimulating factor 2 (granulocyte-macrophage)
 Consensus includes gb:FAW513509 /FEA=EST /DB_XREF=gi:7151587 /DB_XREF=est:xd6d08.x1 /CLONE=IMAGE:2707023 /UG=Hs.254679 ESTs, Highly similar to Y539_H1 AW513509
 TRIAD3 protein
 Consensus includes gb:A659005 /FEA=EST /DB_XREF=gi:4782575 /DB_XREF=est:u23602.x1 /CLONE=IMAGE:2251698 /UG=Hs.4082 lectin, galactoside-binding, soluble, t A659005
 hypothetical protein
 progesterone receptor membrane component 2
 glutamate receptor, ionotropic, kainate 5
 a disintegrin and metalloproteinase domain 28
 LIM domain binding 1
 mannosidase, alpha, class IB, member 1
 hypothetical protein FLJ20847
 acyl-Coenzyme A oxidase 1, palmitoyl
 hypothetical protein FLJ13693
 splicing factor, arginine/serine-rich 2
 melanoma antigen, family B, 3
 vacuolar related kinase 2
 lamin B receptor
 Blauval D homolog 1 (Drosophila)
 pleomorphic adenoma gene-like 1
 similar to S. cerevisiae RER1
 placental protein 13-like protein
 KIAA0406 gene product
 eductin 3 (gamma)
 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1 (7.5kD, MWFE)
 RAOS2 homolog (S. cerevisiae)
 hypothetical protein FLJ22824
 endothelial cell growth factor 1 (platelet-derived)
 prostatic dehydrogenase (lipamide) alpha 1
 nuclear factor I/C (CCAAT-binding transcription factor)
 nucleobindin 1
 secreted and transmembrane 1
 protein phosphatase 2A, regulatory subunit B' (PR 53)
 3'-phosphoadenosine 5'-phosphosulfate synthase 2
 AW299958

hypothetical protein FLJ11467
 refined dehydrogenase homolog
 nuclear protein
 MADS box transcription enhancer factor 2, polypeptide A (myocyte enhancer factor 2A)
 down-regulated in metastasis
 high-mobility group (nonhistone chromosomal) protein 1
 claudin 11 (oligodendrocyte transmembrane protein)
 cortistatin
 KIAA1232 protein
 mucopolin 1
 phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 duncie homolog, Drosophila)
 putative Rab5-interacting protein
 PCTAIRE protein kinase 2
 tropomodulin
 Down syndrome critical region gene 3
 scd-lite 2 (Drosophila)
 KIAA0912 protein
 Consensus includes gb:AL049842/DEF=Human DNA sequence from clone 129L7 on chromosome 6q12-13. Contains the gene for a PUTATIVE novel protein. ESTs, an STS, 'AL049842
 surfactant, pulmonary-associated protein A2
 endothelin receptor type A
 fatty acid desaturase 1
 N-ethylmaleimide-sensitive factor attachment protein, gamma
 Consensus includes gb:AL109714.1/DEF=Homo sapiens mRNA (full length insert cDNA clone EUROMIMAGE 327506. /FEA=mRNA/DB_XREF=gi:5669826 /UG=Hs.195484 Ho
 KIAA0618 gene product
 Consensus includes gb:BF436315 /FEA=EST /DB_XREF=gi:11446630 /DB_XREF=est:7p06b05.x1 /CLONE=IMAGE:3644888 /UG=Hs.296338 ESTs
 adaptor-related protein complex 1, sigma 2 subunit
 sodium channel, voltage-gated, type IX, alpha polypeptide
 vitamin D (1,25-dihydroxyvitamin D3) receptor
 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 4
 retinal G protein coupled receptor
 retinoic acid repressible protein
 neuropeptide FF-amide peptide precursor
 lens epithelial protein
 fructose-1,6-bisphosphatase 1
 endothelin receptor type A
 Consensus includes gb:AA143765 /FEA=EST /DB_XREF=gi:1713188 /DB_XREF=est:z031e02.s1 /CLONE=IMAGE:588506 /UG=Hs.71779 Homo sapiens DNA from chromosome
 CGL-06 protein
 proteasome (prosome, macropain) subunit, beta type, 5
 translocase of inner mitochondrial membrane 8 homolog B (yeast)
 CD8 antigen, beta polypeptide 1 (p27)
 KIAA0193 gene product
 hypothetical protein FLJ2222
 transforming growth factor, beta receptor II (70-80kD)
 Consensus includes gb:AU146945 /FEA=EST /DB_XREF=gi:11008466 /DB_XREF=est:AU146945 /CLONE=HMEMB1001944 /UG=Hs.296667 Homo sapiens cDNA FLJ12040
 aldehyde dehydrogenase 5 family, member A1 (succinate-semialdehyde dehydrogenase)
 KIAA1096 protein
 Consensus includes gb:NA48361 /FEA=EST /DB_XREF=gi:1189527 /DB_XREF=est:yy79g10.s1 /CLONE=IMAGE:279810 /UG=Hs.15787 Homo sapiens mRNA: cDNA DKFZp5
 ATP-D-glucose 4-6-dehydratase
 ADP-ribosylation factor domain protein 1, 64kD
 hypothetical protein MGC3222
 TRIAD3 protein
 actin cross-linking factor
 transformin receptor 2
 Consensus includes gb:AL135960 /DEF=Human DNA sequence from clone RP1-18D14 on chromosome 1 Contains genes for SIL (TAL1 (SCL) interrupting locus), TAL1 (T-ce
 hypothetical protein FLJ20171
 myosin IC
 thymodulin reductase beta
 cell division cycle 34
 malignancy-associated protein
 linker for activation of T cells
 RecQ protein-like 5
 splicing factor, arginine/serine-rich 10 (transformer 2 homolog, Drosophila)
 NM_024863 1.09
 NM_005771 1.09
 AU357871 1.09
 NM_005587 1.09
 AF072718 1.09
 NM_002128 1.09
 NM_005602 1.09
 AL577583 1.09
 AI916274 1.09
 NM_020533 1.09
 U18088 1.09
 NM_018840 1.09
 AF742210 1.09
 AI583383 1.09
 NM_006052 1.09
 BG265616 1.09
 AB020719 1.09
 NM_006926 1.09
 AU118882 1.09
 BG165833 1.09
 BC001889 1.09
 NM_014833 1.09
 BF436315 1.09
 NM_003916 1.09
 NM_002977 1.09
 AA172285 1.09
 NM_004554 1.09
 NM_002921 1.09
 AW188134 1.09
 NM_003717 1.09
 NM_018655 1.09
 O26054 1.09
 S81545 1.09
 NM_001424 1.09
 AA143765 1.09
 NM_015937 1.09
 BC704146 1.09
 NM_012459 1.09
 AW296309 1.09
 NM_014738 1.09
 AI855411 1.09
 NM_003242 1.08
 AU146945 1.08
 NM_001080 1.08
 AL096857 1.08
 NM_014305 1.08
 AF230398 1.08
 W74580 1.08
 NM_019011 1.08
 AK023406 1.08
 NM_003227 1.08
 NM_013590 1.08
 NM_017697 1.08
 X96507 1.08
 AF201385 1.08
 BG476681 1.08
 AF041410 1.08
 AF038905 1.08
 AB042823 1.08
 NM_004593 1.08

Cystatin EM	NM_001323	1.08
leucine zipper, putative tumor suppressor 1	AF123856	1.08
Splicing factor, arginine/serine-rich, 48KD	A1343249	1.08
tumor endothelial marker 7 precursor	NM_020405	1.08
acrosomal vesicle protein 1	NM_020115	1.08
hypothetical protein FLJ10314	NM_018055	1.08
hypothetical protein FLJ20272	NM_017735	1.08
hypothetical protein MGC2828	NM_024076	1.08
cubilin (intrinsic factor-cobalamin receptor)	NM_001081	1.08
SKIP for skeletal muscle and kidney enriched inositol phosphatase	NM_016532	1.08
immunoglobulin mu binding protein 2	L14754	1.08
tumor-associated calcium signal transducer 2	J04152	1.08
KIAA0349 protein	A1761518	1.08
sequestrinome 1	NM_003900	1.08
KIAA0841 protein	AB020646	1.08
myeloid leukemia factor 1	NM_022443	1.08
poly(A) binding protein, nuclear 1	A1130920	1.08
guanylate cyclase activator 2B (uroguanylin)	NM_007102	1.08
cisplatin resistance-associated overexpressed protein	NM_016424	1.08
neuropeptide Y receptor Y2	NM_000910	1.08
mannose-6-phosphate receptor (calnexin dependent)	NM_002355	1.08
KIAA1067 protein	BE646386	1.08
nuclear factor related to kappa B binding protein	NM_006165	1.08
heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37KD)	D55672	1.08
MAP/microtubule affinity-regulating kinase like 1	AB049127	1.08
CDC7 cell division cycle 7-like 1 (S. cerevisiae)	NM_017717	1.08
PTD015 protein	NM_003503	1.08
solute carrier family 15 (H ⁺ /peptide transporter), member 2	BC002752	1.08
collectin sub-family member 12	BF223679	1.08
fibroblast growth factor 22	NM_030781	1.08
four and a half LIM domains 1	NM_020637	1.08
DKFZP566J0619 protein	NM_001449	1.08
kinesin heavy chain member 2	BF055496	1.08
glycogenin	BE872563	1.08
topoisomerase (DNA) II alpha (170KD)	AF087842	1.08
chromosome 21 open reading frame 91	AL581834	1.08
testis specific ankyrin-like protein 1	NM_017447	1.08
KIAA0923 protein	NM_018552	1.08
hypothetical protein FLJ20967	AK001710	1.08
collagen, type VI, alpha 2	NM_022071	1.08
CD1D antigen, d polypeptide	AY029208	1.08
ubiquitin-cytochrome c reductase binding protein	NM_001766	1.08
KIAA0176 protein	BC005230	1.08
Consensus includes gb:AW006893 /FEA=EST /DB_XREF=gi:5855731 /DB_XREF=est:w08606.x1 /CLONE=IMAGE:2506859 /UG=Hs.237946 ESTs, Moderately similar to ALL	A4523921	1.08
imidazole receptor candidate	NM_007184	1.08
hypothetical protein FLJ12118	NM_024537	1.08
hemochromatosis	AF148804	1.08
hypothetical protein FLJ14069	NM_025023	1.08
DKFZP564G196 protein	AL049329	1.08
apolipoprotein A-I	NM_000039	1.08
hypothetical protein FLJ20207	NM_017711	1.08
myosin, light polypeptide 9, regulatory	NM_006097	1.08
radixin	AL137251	1.08
replication factor C (activator 1) 5 (36.5KD)	BC001868	1.08
regulator of G-protein signalling 12	AA687475	1.08
Consensus includes gb:AL046979 /FEA=EST /DB_XREF=gi:5435035 /DB_XREF=est:DKFZp568K0617 /UG=Hs.9973 tensin	AL046979	1.08
Consensus includes gb:AL574186 /FEA=EST /DB_XREF=gi:12934150 /DB_XREF=est:AL574186 /CLONE=CS001039YM09 (3 prime) /UG=Hs.9739 glycerol-3-phosphate dehyd	AL574186	1.08
trichocoid repeat containing 11 (THR-associated protein, 230 kDa subunit)	AF132033	1.08
Consensus includes gb:NT3931 /FEA=EST /DB_XREF=gi:1231216 /DB_XREF=est:z674a07.s1 /CLONE=IMAGE:298260 /UG=Hs.323079 Homo sapiens mRNA; cDNA DKFZp	NT3931	1.08
KIAA0818 protein	NM_015057	1.08
complement component 1, q subcomponent binding protein	M69039	1.08
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 15	NM_001358	1.08

tumor necrosis factor receptor superfamily, member 6	AA164751	1.08
KIAA1448 protein	AI06628	1.08
plasminogen activator, urokinase receptor	AC005525	1.08
TATA box binding protein	NM_003194	1.08
serpin trypsin-like protease	NM_004262	1.08
zinc finger protein 204	AL021808	1.08
HIV-1 Tat interactive protein, 60 kDa	NM_006368	1.08
hypothetical protein FLJ20199	AK00253	1.08
calcium/calmodulin-dependent serine protein kinase (MAGUK family)	NM_003668	1.08
butyrophilin, subfamily 2, member A1	NM_007049	1.08
cholinergic receptor, muscarinic 4	NM_000741	1.08
butyrophilin, subfamily 2, member A1	U90142	1.08
matrin 3	AA129420	1.08
hypothetical protein FLJ10052	NM_017882	1.08
E4F transcription factor 1	NM_004424	1.08
hypothetical protein FLJ2127	NM_022775	1.08
homoeo box A7	AF026397	1.08
protein kinase, X-linked	NM_005044	1.08
BCL2-like 1	U72386	1.08
stil homolog 1 (Drosophila)	AB011537	1.08
ubiquitin-conjugating enzyme E2H (UBC8 homolog, yeast)	AB239520	1.08
a disintegrin and metalloproteinase domain 9 (mellin gamma)	NM_003816	1.08
KIAA0036 gene product	NM_014642	1.08
neural killer-tumor recognition sequence	NM_005385	1.08
hypothetical protein MGC11335	NM_030819	1.08
3-hydroxy-3-methylglutaryl-Coenzyme A synthase 1 (soluble)	BG035985	1.08
paired box gene 9	NM_006194	1.08
hypothetical protein FLJ23380	NM_023076	1.08
Rev/Rax activation domain binding protein-related	U727494	1.08
signal recognition particle 19kD	NM_003135	1.08
KIAA0999 protein	A4044154	1.08
UDP-GlcNAc 6-phosphatase catalytic subunit-related protein	D29805	1.08
KIAA0795 protein	NM_021176	1.08
fatty acid binding protein 7, brain	AB018338	1.08
branched chain keto acid dehydrogenase E1, alpha polypeptide (maple syrup urine disease)	NM_001446	1.08
KIAA1453 protein	NM_000709	1.08
Consensus includes gb-AL162044.1 /DEF=Homo sapiens mRNA, cDNA DKFZ761L0812 (from clone DKFZ761L0812), partial cds. /FEA=mRNA /GEN=DKFZ761L0812 /PF AL162044	NM_025090	1.08
collagen, type IV, alpha 2	AA909035	1.08
ADP-ribosylation factor-like 4	NM_005738	1.08
integrin, alpha 7	AK022548	1.08
DC-specific transmembrane protein	NM_030788	1.08
hypothetical protein FLJ14393	AB23896	1.08
protein phosphatase 1, regulatory (inhibitor) subunit 16B	AB020630	1.08
hypothetical protein FLJ20397	NM_017802	1.08
proline synthetase co-transcribed (bacterial homolog)	AA176833	1.08
Consensus includes gb-AK025343.1 /DEF=Homo sapiens cDNA, FLJ21690 fls, clone COL09538. /FEA=mRNA /DB_XREF=gj:10437841 /UG=Hs.306802 Homo sapiens cDNA: AK025343	NM_018226	1.08
arginyl aminopeptidase (aminopeptidase B)-like 1	NM_018141	1.08
mitochondrial ribosomal protein S10	NM_02480	1.08
protein phosphatase 1, regulatory (inhibitor) subunit 12A	AF070448	1.08
cathepsin L2	BC002600	1.08
mature T-cell proliferation 1	NM_014423	1.08
KIAA0368 protein	AB002366	1.08
cholinergic receptor, nicotinic, beta polypeptide 1 (muscle)	NM_000747	1.08
Bart-Hill homeobox 1	NM_021570	1.08
hypothetical protein FLJ22191	NM_025231	1.08
SP140 nuclear body protein	NM_007237	1.08
ADP-ribosylation factor 6	A67763	1.08
trinucleotide repeat containing 5	BC004423	1.08
KIAA0332 protein	AB002330	1.08
FAST kinase	NM_006712	1.08
thioredoxin-like 2	AK022131	1.08

brain and reproductive organ-expressed (TNFRSF1A modulator)	NM_004899	1.08
SEC23 protein	NM_007214	1.08
DKFZP586C1820 protein	AV694093	1.08
casein kinase 1, alpha 1	BF310713	1.08
nuclear receptor co-repressor 2	AI498095	1.08
succinate dehydrogenase complex, subunit D, integral membrane protein	NM_003002	1.08
insulin-like growth factor binding protein 7	NM_001553	1.08
Rab geranylgeranyltransferase, beta subunit	U49245	1.08
zinc finger protein 181 (pZ-67)	NM_003443	1.08
RAP1A, member of RAS oncogene family	NM_002884	1.08
hypothetical protein FLJ10276	NM_018045	1.08
LATS, large tumor suppressor, homolog 1 (Drosophila)	NM_004690	1.08
KIAA0481 gene product	T51252	1.08
hypothetical protein FLJ22679	NM_017698	1.08
nemo-like kinase	NM_016231	1.08
complement component 1, q subcomponent, beta polypeptide	NM_000491	1.08
XPA binding protein 1; putative ATP[GTP]-binding protein	AB044661	1.08
ferrochialase (protophyria)	AU152635	1.08
patent protein 13	NM_013268	1.08
hemopexin	M35803	1.08
suppressor of variegation 3-9 homolog 1 (Drosophila)	NM_003173	1.08
synthase 8	AF130102	1.08
retinoic acid repressible protein	NM_004853	1.08
Friedreich ataxia	NM_000144	1.08
breast cancer 2, early onset	X95152	1.08
glutamate dehydrogenase 1	AJ339331	1.08
hypothetical protein FLJ12584	NM_025139	1.08
pyruvate dehydrogenase (lipoamide) beta	AL17618	1.08
butyrophilin, subfamily 3, member A1	U90552	1.08
STRIN protein	NM_016271	1.08
DKFZP586I2223 protein	AL080214	1.08
nephroblastoma overexpressed gene	NM_002514	1.08
dynlen, axonemal, light polypeptide 4	NM_005740	1.08
CASP8 and FADD-like apoptosis regulator	AF009619	1.08
Consensus includes gb:AA744682 FEA=EST		1.08
testis expressed sequence 14	NM_031272	1.08
UDP glycosyltransferase 1 family, polypeptide A9	NM_021027	1.08
KIAA0489 protein	AB007958	1.08
hypothetical protein FLJ20457	NM_017832	1.08
protein tyrosine phosphatase, receptor type, C	AI809341	1.08
mammian-binding lectin serine protease 1 (C4/C2 activating component of Rar-reactive factor)	D17525	1.08
ceroid-lipofuscinosis, neuronal 2, late infantile (Jensky-Bielschowsky disease)	NM_000391	1.08
HCF-binding transcription factor Zhangfei	AI432137	1.08
erlavin 1	AL044651	1.08
ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1	AV711183	1.08
mitochondrial intermediate peptidase	NM_005932	1.08
clustrin (complement lysin inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	M25915	1.08
hypothetical protein FLJ10751	NM_018205	1.07
flucodrase, alpha-L-1, tissue	NM_000147	1.07
SFRS protein kinase 1	NM_003137	1.07
cholinergic receptor, nicotinic, gamma polypeptide	NM_005199	1.07
actin related protein 2/3 complex, subunit 4 (20 kD)	NM_005718	1.07
cell adhesion molecule-related/down-regulated by oncogenes	NM_016952	1.07
complement C11-like proteinase precursor,	NM_016546	1.07
arylsulfatase D	NM_001669	1.07
hypothetical protein FLJ22160	NM_024585	1.07
mitogen-activated protein kinase 7	NM_002749	1.07
dystonia 1, torsion (autosomal dominant; torsin A)	NM_000113	1.07
glutamate receptor, metabotropic 8	AC000099	1.07
hypothetical protein FLJ12879	NM_024757	1.07
hypothetical protein FLJ23469	NM_024710	1.07
HSPC040 protein	NM_016098	1.07
protein tyrosine phosphatase type IVA, member 1	AW165960	1.07

sequestosome 1 1.07 N30649

neural precursor cell expressed, developmentally down-regulated 4 1.07 D42055

chromosome 1 open reading frame 2 1.07 NM_006569

tumor suppressing subtransferable candidate 1 1.07 NM_003310

Mid-1-related chloride channel 1 1.07 AW195581

lec protein brosinase kinase 1.07 NM_003215

5,10-methylenetetrahydrofolate reductase (NADPH) 1.07 NM_003957

G protein-coupled receptor 50 1.07 NM_004224

HeLa cyclin-dependent kinase 2 interacting protein 1.07 NM_016550

actin related protein 2/3 complex, subunit 2 (34 kD) 1.07 NM_005731

A kinase (PRKA) anchor protein 11 1.07 NM_016248

intercellular adhesion molecule 1 (ICSA), human rhinovirus receptor 1.07 AA284705

tumor repeat associator with PCTAIRE 2 1.07 AW129593

cytochrome P450, subfamily IIC (methoxytol 4-hydroxylase), polypeptide 9 1.07 NM_000771

KIAA0680 gene product 1.07 NM_014721

DNA1 dosage suppressor of mck1 homolog, meiosis-specific homologous recombination (yeast) 1.07 NM_007068

ornithine decarboxylase anlyzyme 1 1.07 AF090094

UMP-CMP kinase 1.07 NM_016308

24 kDa intrinsic membrane protein 1.07 BF057649

Cytochrome P450, subfamily XXVIB (25-hydroxyvitamin D-1-alpha-hydroxylase), polypeptide 1 1.07 NM_000785

Consensus includes gb:W48843 /FEA=EST /DB_XREF=gi:1337022 /DB_XREF=est:zc44a09.s1 /CLONE=IMAGE:325144 /UG=Hs:285814 sprouty (Drosophila) homolog 4 /FL: W48843 1.07

cytochrome P450, subfamily IIO (debrisoquine, sparteine, etc., -metabolizing), polypeptide 6 1.07 NM_000106

proprotein convertase subtilisin/kexin type 1 inhibitor 1.07 NM_013271

heterogeneous nuclear ribonucleoprotein D-like 1.07 A1762552

NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 1 (7ND, MNLL) 1.07 NM_004545

oxytocin, pramoxine (neurophysin II) 1.07 NM_000915

gb:NM_017421.1 /DEF=Homo sapiens methyltransferase COQ3 (COQ3), mRNA, /FEA=mRNA /GEN=COQ3 /PROD=methyltransferase COQ3 /DB_XREF=gi:8393176 /FL=gb: NM_017421 1.07

ankyrin repeat domain 6 1.07 NM_014942

Probe nt1737 (polycystic kidney disease, autosomal recessive, in) 1.07 NM_006531

ATP binding protein associated with cell differentiation 1.07 NM_005783

serum amyloid A4, constitutive 1.07 NM_006512

Homer, neuronal immediate early gene, 3 1.07 AC002985

dysenin, axonemal, heavy polypeptide 9 1.07 AF257737

Insitol 1,4,5-triphosphate receptor, type 1 1.07 U23850

Consensus includes gb:AL096814 /DEF=Human DNA sequence from clone RP1-139D8 on chromosome 6p12.1-21.1 Contains genes for GUCA1A (guanylate cyclase activator) AL096814 1.07

gb:NM_001222.1 /DEF=Homo sapiens calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma (CAMK2G), mRNA, /FEA=mRNA /GEN=CAMK2G /PROD=calcium NM_001222 1.07

PRO2000 protein 1.07 NM_014109

hypothetical protein FLJ22781 1.07 NM_025130

KIAA0701 protein 1.07 AA05798

actinin, alpha 4 1.07 U48734

active BCR-related gene 1.07 L19704

solute carrier family 5 (sodium-dependent vitamin transporter), member 6 1.07 NM_021095

Consensus includes gb:AU148040 /FEA=EST /DB_XREF=gi:1009561 /DB_XREF=est:AU148040 /CLONE=MAMMA1002428 /UG=Hs:301615 Homo sapiens cDNA FLJ12370 AU148040 1.07

Consensus includes gb:U21915.1 /DEF=Human chromosome 5q13.1 clone 5G8 mRNA, /FEA=mRNA /DB_XREF=gi:736412 /UG=Hs:14658 Human chromosome 5q13.1 clone U21915 1.07

gb:AF274954.1 /DEF=Homo sapiens PNAS-28 mRNA, complete cds, /FEA=mRNA /PROD=PNAS-28 /DB_XREF=gi:12751070 /UG=Hs:326791 Homo sapiens PNAS-28 mRNA /AF274954 1.07

gb:NM_017871.1 /DEF=Homo sapiens hypothetical protein FLJ20542 (FLJ20542), mRNA, /FEA=mRNA /GEN=FLJ20542 /PROD=hypothetical protein FLJ20542 /DB_XREF=gi: NM_017871 1.07

dishevelled, dish homolog 3 (Drosophila) 1.07 NM_004423

KIAA0301 protein 1.07 BE670928

programmed cell death 2 1.07 NM_002588

zinc finger protein 93 homolog (mouse) 1.07 AC084239

activating transcription factor 5 1.07 BC005174

ribosomal protein S13 1.07 NM_001017

Ca2+-promoted Ras inactivator 1.07 A1736591

CAS98 associated protein 2 1.07 AB037736

erythrocyte membrane protein band 4.1 (elliptocytosis 1, RH-linked) 1.07 AF156225

nucleoporin 54kD 1.07 NM_017426

membrane protein of cholinergic synaptic vesicles 1.07 BC001913

hypothetical protein similar to RNA-binding protein Iark 1.07 BC003503

actin, alpha 2, smooth muscle, aorta 1.07 NM_001613

ARPI actin-related protein 1 homolog A, contractin alpha (yeast) 1.07 NM_005736

integrin, beta 6 1.07 NM_002214

serpin/SUMO-specific protease 1.07 NM_020854

glutamate receptor, ionotropic, AMPA 2 1.07 NM_000826

HLA class II region expressed gene KE4
 plasminogen
 KIAA0107 gene product
 Consensus includes gb:BE780075 /FEA=EST /DB_XREF=gi:10201273 /DB_XREF=est:60146954F1 /CLONE=IMAGE:3873157 /UG=Hs.74137 transmembrane trafficking protein BE780075
 HLA-B associated transcript 2
 actn, alpha 1, skeletal muscle
 arachidonate 5-lipoxygenase
 xeroderma pigmentosum, complementation group A
 integrin beta 1 binding protein (melusin) 2
 CAT56 protein
 Consensus includes gb:U00784.1 /DEF=Human EST clone 122887 mariner transposon Hamar1 sequence, /FEA=mRNA /DB_XREF=gi:2231387 /UG=Hs.132967 Human EST
 ATP-binding cassette, sub-family F (GCN20), member 2
 CLIP-associating protein 2
 protein tyrosine phosphatase type IVA, member 2
 Consensus includes gb:AL049280.1 /DEF=Homo sapiens mRNA: cDNA DKFZ6564K143 /FEA=mRNA /DB_XREF=gi:4500037 /UG=Hs.155387 FAL049280
 Consensus includes gb:AF070536.1 /DEF=Homo sapiens clone 24566 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3387893 /UG=Hs.133342 Homo sapiens clone 24566 mF AF070536
 BCL2adenovirus E1B 19kD interacting protein 1
 hypothetical protein FLJ10948
 hypothetical protein MGC3178
 zinc finger protein 10 (KIX 1)
 Consensus includes gb:AJ249275 /DEF=Homo sapiens partial MTHFR gene for methyltetrahydrofolate reductase /FEA=CDS /DB_XREF=gi:7271187 /UG=Hs.214142 S.10-AJ249275
 dimethylarginine dimethylaminohydrolase 2
 mitogen-activated protein kinase Kinase 2
 major intrinsic protein of lens fiber
 spectrin, beta, non-erythrocytic 4
 sulfolipase family, cytosolic, 1A, phenol-preferring, member 1
 hypothetical protein FLJ22439
 chromosome 1 open reading frame 13
 heterogeneous nuclear ribonucleoprotein L
 recoverin
 CTD (carboxy-terminal domain, RNA polymerase II, polypeptide A) phosphatase, subunit 1
 comedodermosin
 hypothetical protein MGC2494
 thrombopoietin (myeloproliferative leukemia virus oncogene ligand, megakaryocyte growth and development factor)
 olfactory receptor, family 2, subfamily B, member 6
 surfactant, pulmonary-associated protein C
 KIAA1068 protein
 lucosyltransferase 7 (alpha (1,3) lucosyltransferase)
 bactericidal/permeability-increasing protein
 hexose-6-phosphate dehydrogenase (glucose 1-dehydrogenase)
 gb:BC002877.1 /DEF=Homo sapiens. Similar to hypothetical protein FLJ11585, clone MGC:11258, mRNA complete cds, /FEA=mRNA /PROD=Similar to hypothetical protein
 PTPRF interacting protein, binding protein 1 (liprin beta 1)
 immunoglobulin kappa constant
 KIAA0453 protein
 KIAA0543 protein
 Consensus includes gb:AF052103.1 /DEF=Homo sapiens clone 23726 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3360410 /UG=Hs.159157 Homo sapiens clone 23726 mF AF052103
 phosphatidylinositol-4-phosphate 5-kinase, type I, alpha
 eta variant gene 5 (eta-related molecule)
 myogenic factor 6 (herculin)
 hypothetical protein MGC10731
 RAD17 homolog (S. pombe)
 immunoglobulin lambda locus
 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 3
 protein tyrosine phosphatase, non-receptor type 13 (APO-1/CD95, Fas)-associated phosphatase)
 nucleosome assembly protein 1-like 1
 hypothetical protein MGC2663
 gogli autoantigen, pcdgn subfamily b, macrophage (with transmembrane signal), 1
 farnesyl-diphosphate farnesyltransferase 1
 Consensus includes gb:AK022331.1 /DEF=Homo sapiens cDNA FLJ12269 (fa, clone MAMMA1001635, /FEA=mRNA /DB_XREF=gi:10433706 /UG=Hs.293925 Homo sapiens
 iduronidase, alpha-L-
 SWAP-70 protein
 Consensus includes gb:A638420 /FEA=EST /DB_XREF=gi:4690654 /DB_XREF=est:131g07.x1 /CLONE=IMAGE:2242428 /UG=Hs.25035 chloride intracellular channel 4
 cathecin 8, type 2

TIAT1 cytoplasmic granule-associated RNA binding protein	AL046419	1.06
cyclochrome P450, subfamily 1D (debrisoquine, sparteine, etc., -metabolizing), polypeptide 6	X18866	1.06
hypothetical protein FL20320	NM_017765	1.06
serding nexin 10	NM_010322	1.06
hypothetical protein FLJ23529	AK027182	1.06
ribosomal protein L37	BF216701	1.06
Consensus includes gb:AL023773 /DEF=Human DNA sequence from clone 406C18 on chromosome Xq27.1-27.3. Contains two MAGE (Melanoma Associated Antigen) family	AL023773	1.06
HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae)	AL570294	1.06
KUAA0759 protein	AB018302	1.06
zinc finger protein 283	NM_005741	1.06
nesca protein	NM_014328	1.06
KUAA0657 protein	AI633851	1.06
odorant-binding protein 2A	NM_014582	1.06
ring finger protein 10	AL578551	1.06
NgpV17 transgene, murine homolog, glomerulosclerosis	NM_002437	1.06
KUAA0441 gene product	NM_014797	1.06
Consensus includes gb:AL137673.1 /DEF=Homo sapiens mRNA: cDNA DKFZp343H0872 (from clone DKFZp343H0872). /FEA=mRNA /DB_XREF=gi:16607841 /UG=Hs.30645	AL137673	1.06
Consensus includes gb:AW972351 /FEA=EST /DB_XREF=gi:8162197 /DB_XREF=est:EST384442 /UG=Hs.293451 ESTs	AW972351	1.06
hypothetical protein FLJ10713	NM_018189	1.06
Consensus includes gb:L43578.1 /DEF=Homo sapiens (clone 115392) mRNA. /FEA=mRNA /DB_XREF=gi:1895067 /UG=Hs.82171 Homo sapiens (clone 115392) mRNA	L43578	1.06
mitogen inducible 2	AW469573	1.06
hypothetical protein FLJ22531	NM_024650	1.06
MHC class II transactivator	NM_000246	1.06
zinc finger protein 226	NM_015819	1.06
shu1n silent mating type information regulation 2 homolog 6 (S. cerevisiae)	NM_016539	1.06
mitogen-activated protein kinase kinase 5	U71087	1.06
SART1 protein	AY008268	1.06
TIAT1 cytoplasmic granule-associated RNA binding protein	H85548	1.06
hypothetical protein FLJ12598	NM_024754	1.06
Consensus includes gb:Z95624 /DEF=Human DNA sequence from cosmid U237H1 contains Ras like GTPase and ESTs /FEA=CDS /DB_XREF=gi:2117165 /UG=Hs.248013 P_Z95624	NM_024754	1.06
Consensus includes gb:BF882927 /FEA=EST /DB_XREF=gi:12385739 /DB_XREF=est:502306318F1 /CLONE=IMAGE:4397525 /UG=Hs.324787 soluble carrier family 5 (mosillo BF882927	BF882927	1.06
Consensus includes gb:N36926 /FEA=EST /DB_XREF=gi:1158068 /DB_XREF=est:y39e04.s1 /CLONE=IMAGE:273534 /UG=Hs.286361 potassium inwardly-rectifying channel N36926	N36926	1.06
carboxypeptidase M	AV710357	1.06
T-box 8	NM_004608	1.06
G protein-coupled receptor 63	NM_030784	1.06
protein kinase C, theta	L01087	1.06
HSV associated factor	NM_006462	1.06
ribosomal protein L27	NM_000968	1.06
chromosome 17 open reading frame 31	AB018275	1.06
KUAA0008 gene product	NM_014750	1.06
heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1	NM_006041	1.06
hypothetical protein MGC2744	AI083187	1.06
KUAA0943 protein	AL080168	1.06
joined to JAZF1	BF382924	1.06
fragile X mental retardation, autosomal homolog 2	NM_004860	1.06
cytochrome c	BC052999	1.06
candidate tumor suppressor p33 ING1 homolog	NM_016162	1.06
protein phosphatase 2 (formerly 2A), catalytic subunit, alpha isoform	BC000400	1.06
hyaluronan synthase 2	NM_005328	1.06
Smr2-related CBP activator protein	AB002207	1.06
BENE protein	BC003179	1.06
hypothetical protein FLJ20138	NM_017684	1.06
v-maf musculosarcomatous fibrosarcoma oncogene homolog (avian)	AF055376	1.06
hypothetical protein FLJ12729	NM_024847	1.06
Consensus includes gb:A050783 /FEA=EST /DB_XREF=gi:3307598 /DB_XREF=est:ov12509.x1 /CLONE=IMAGE:1637081 /UG=Hs.125064 Homo sapiens chromosome 19, c A050783	NM_024847	1.06
DEADH (Asp-Glu-Ala-Asp-His) box binding protein 1	BC004331	1.06
GAP-associated tyrosine phosphoprotein p62 (Sam68)	N24668	1.06
reproduction 6	NM_006559	1.06
optic atrophy 3 (autosomal recessive, with chorea and spastic paraplegia)	O63768	1.06
catechol-O-methyltransferase	NM_025136	1.06
ceroid-lipofuscinosis, neuronal 8 (epilepsy, progressive with mental retardation)	BC000419	1.06
hypothetical protein MGCA174	NM_018841	1.06
	NM_024319	1.06

DXFZP564G092 protein
 spinocerebellar ataxia 1 (olivopontocerebellar ataxia 1, autosomal dominant, ataxin 1)
 hypothetical protein FLJ22871
 KIAA1058 protein
 KIAA0101 gene product
 hypothetical protein FLJ10914
 cullin 3
 diphtheria toxin resistance protein required for diphtheramide biosynthesis (Saccharomyces)-like 2
 megakaryoblastic leukemia (translocation) 1
 ATP-binding cassette, sub-family B (MDR/TAP), member 1
 hypothetical protein FLJ13087
 neuronal precursor cell expressed, developmentally down-regulated 4-like
 PRO1648 protein
 Consensus includes gb:BG187570 /FEA=EST /DB_XREF=gi:12674199 /DB_XREF=est:602342795F1 /CLONE=IMAGE:4453055 /UG=Hs.321717 ESTs, Weakly similar to S22: BG167570 X03348
 nuclear receptor subfamily 3, group C, member 1
 UDP-N-acetyl-alpha-D-glucosamine polypeptide N-acetylglucosaminyltransferase 1 (GalnAc-T1)
 osteomodulin
 filamin A, alpha (ectin binding protein 280)
 prolyl endopeptidase
 nuclear receptor subfamily 4, group A, member 3
 Nef-associated factor 1
 hypothetical protein MGC5468
 inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase gamma
 frizzled homolog 1 (Drosophila)
 clusterin (complement lysis inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)
 dUTP pyrophosphatase
 Consensus includes gb:AC005756 /DEF=Homo sapiens chromosome 19, fosmid 39347 /FEA=CDS /DB_XREF=gi:3688091 /UG=Hs.247754 Homo sapiens chromosome 19, ic AC005756 U62891
 melanoma antigen, family D, 2
 DXFZP34A043 protein
 ribosomal protein S4, Y-linked
 Ras protein-specific guanine nucleotide-releasing factor 1
 ras homolog gene family, member F (in filopodia)
 pellino homolog 1 (Drosophila)
 galactokinase 1
 hypothetical protein FLJ21034
 myoidenium cofactor sulfatase
 hypothetical protein FLJ20315
 aminomethyltransferase (glycine cleavage system protein T)
 bone morphogenetic protein 3 (osteogenic)
 interleukin 9 receptor
 Consensus includes gb:BC235929 /FEA=EST /DB_XREF=gi:12749776 /DB_XREF=est:naf18n02.x1 /CLONE=IMAGE:4141658 /UG=Hs.26034 ESTs
 Consensus includes gb:AK024673.1 /DEF=Homo sapiens cDNA: FLJ21026 fls, clone CAE06812, /FEA=mRNA /DB_XREF=gi:10437019 /UG=Hs.317585 Homo sapiens cDNA: AK024679
 Consensus includes gb:AW166925 /DB_XREF=gi:16398450 /DB_XREF=est:tg88601.x1 /CLONE=IMAGE:2633456 /UG=Hs.289037 Homo sapiens cDNA FLJ14135 AW166925
 karyopherin alpha 1 (importin alpha 5)
 KIAA1078 protein
 diphtheria toxin receptor (heparin-binding epidermal growth factor-like growth factor)
 chromosome 12 open reading frame 2
 carbonyl triate (chondroitin) synthase 1
 heparin-binding protein
 Clusar incl. A4392932:mb0a11.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:1090172 /clone_end=3 /gb=AA582932 /gi=2360292 /ug=Hs.21349 /len=614
 GTP-binding protein homologous to Saccharomyces cerevisiae SEC4
 immunoglobulin kappa constant
 Consensus includes gb:AB050195.1 /DEF=Homo sapiens mRNA, partial cds, clone:H2-5, MJD homolog, /FEA=mRNA /DB_XREF=gi:11559487 /UG=Hs.302079 Homo sapiens cDNA: AB050195
 Consensus includes gb:S76346.1 /DEF=AML1-1/AML1 (alternatively spliced, exons 5 and 6) human, mRNA, /FEA=mRNA /GEN=AML1 /PROD=AML1 /DB_XREF=S76346
 Consensus includes gb:BC005066.1 /DEF=Homo sapiens, clone IMAGE:3347954, mRNA, partial cds, /FEA=mRNA /PROD=Unknown (protein for IMAGE:3347954) /DB_XREF=BC005066
 early endosome antigen 1, 162KD
 L3424 B subtilis dapB, jpfF, jpfG genes corresponding to nucleolides 1359-3197 of L38424 (-S, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
 KIAA0561 protein
 BAI1-associated protein 3
 hypothetical protein FLJ23305
 Consensus includes gb:R87507 /FEA=EST /DB_XREF=gi:946320 /DB_XREF=est:ym91006.s1 /CLONE=IMAGE:166259 /UG=Hs.12509 Homo sapiens clone 23583 mRNA sec R87507
 forward box E3
 interferon regulatory factor 4

adaptor-related protein complex 4, sigma 1 subunit
 Consensus includes gb:AL353948.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761P1114 (from clone DKFZp761P1114). /FEA=mRNA /DB_XREF=gi:7669990 /UG=Hs.306500
 ribosomal protein S21
 goli autanigen, golgin subfamily a, 2
 Consensus includes gb:AL139377 /DEF=Human DNA sequence from clone RP11-25 J8 on chromosome 13 Contains ESTs, STSs, GSSs and a CpG island. Contains two new
 hypothetical protein FLJ22294
 integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41B)
 brain and reproductive organ-expressed (TNFRSF1A modulator)
 A kinase (PRKA) anchor protein 3
 adenomatous polyposis cell like
 neuroblastoma, suppression of tumorigenicity 1
 biglycan
 ATP-binding cassette, sub-family A (ABC1), member 2
 collagen, type IV, alpha 3 (Goodpasture antigen)
 apolipoprotein C-II
 Consensus includes gb:AU147983 /FEA=EST /DB_XREF=gi:11009504 /DB_XREF=est:AU147983 /CLONE=MAMMA1002308 /UG=Hs.296716 Homo sapiens cDNA FLJ12349
 proto-oncogene bcr-abl
 hypothetical protein FLJ23471
 myo-histidin 1-phosphate synthase A1
 N-myc downstream-regulated gene 2
 GAP-associated tyrosine phosphoprotein p62 (Sam68)
 autocrine motility factor receptor
 KIAA0701 protein
 Fxr1 protein
 hypothetical protein FLJ13788
 thymidine kinase 2, mitochondrial
 density-regulated protein
 KIAA0860 protein
 KIAA0852 gene product
 hypothetical protein FLJ10498
 CG1-94 protein
 ADP-ribosylation factor domain protein 1, 64kD
 hypothetical protein FLJ20116
 KIAA1185 protein
 KIAA1649 protein
 latent transforming growth factor beta binding protein 1
 heterogeneous nuclear ribonucleoprotein D-like
 calcium modulating ligand
 chorionic somatomammotropin hormone 1 (placental lactogen)
 SPTF-associated factor 65 gamma
 core-binding factor, runt domain, alpha subunit 2, translocated to, 1; cyclin D-related
 regulator of G-protein signalling 11
 KIAA0285 protein
 HIC1-related gene on chromosome 22
 potassium voltage-gated channel, shaker-related subfamily, beta member 1
 mitochondrial ribosomal protein 63
 Consensus includes gb:AK022065.1 /DEF=Homo sapiens cDNA FLJ12003 fts, clone HEMBB1001537. /FEA=mRNA /DB_XREF=gi:10433385 /UG=Hs.296683 Homo sapiens c
 hypothetical protein FLJ12571
 GPI-anchored metastase-associated protein homolog
 BA11-associated protein 2
 glutamate receptor, ionotropic, N-methyl D-aspartate 1
 KIAA1054 protein
 KIAA1053 protein
 cold shock domain protein A
 ATPase, H⁺-transporting, lysosomal (vacuolar proton pump) non-catalytic accessory protein 1A (110/118kD)
 protein kinase, lysine deficient 1
 intersectin 2
 hypothetical protein MGC17330
 translocating chain-associating membrane protein
 chloride intracellular channel 4
 protein kinase C, epsilon

protein phosphatase 2, regulatory subunit B (B56), delta isoform
 UDP-N-acetyl-alpha-D-galactosamine:polypptide N-acetylglucosaminyltransferase 1 (GalNAc-T1)
 RAB14, member RAS oncogene family
 androgen receptor (dihydrotestosterone receptor; testicular feminization; spinal and bulbar muscular atrophy; Kennedy disease)
 Consensus includes gb:AL534702 /FEA=EST /DB_XREF=gi:12798195 /DB_XREF=est:AL534702 /CLONE=CSODP0067N17 (3 prime) /UG=Hs.13377 Homo sapiens clone 23c AL534702
 hypothetical protein FLJ10847
 polp1-87
 uncoupling protein 2 (mitochondrial, proton carrier)
 Consensus includes gb:AU146580 /FEA=EST /DB_XREF=gi:11008101 /DB_XREF=est:AU146580 /CLONE=HEMBB1000893 /UG=Hs.296677 Homo sapiens cDNA FLJ11956
 hypothetical protein FLJ20334
 CamKII-like protein kinase
 butyrophilin, subfamily 2, member A1
 sorting nexin 13
 CGI-82 protein
 immediate early response 3
 tracheal collagen repeat containing 11 (THR-associated protein, 230 kDa subunit)
 CD8 antigen
 Consensus includes gb:BE299495 /FEA=EST /DB_XREF=gi:183243 /DB_XREF=est:60094774T1 /CLONE=IMAGE:2960610 /UG=Hs.218329 hypothetical protein /FL=gb:NM
 ATPase inhibitor precursor
 RAI1GDS-like gene
 Consensus includes gb:BF185104 /FEA=EST /DB_XREF=gi:11081827 /DB_XREF=est:7n15a07.x1 /CLONE=IMAGE:3584565 /UG=Hs.56729 lymphocyte-specific protein 1
 Cluster incl. A1524138:th09103.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2117789 /clone_end=3 /gb=AL524138 /gi=4438273 /ug=Hs.23391 len=834
 cAMP responsive element binding protein-like 1
 DKFZP566C0424 protein
 Consensus includes gb:AU147830 /FEA=EST /DB_XREF=gi:11009351 /DB_XREF=est:AU147830 /CLONE=MAMMA1001818 /UG=Hs.296713 Homo sapiens cDNA FLJ12285
 ankyrin 1, erythrocytic
 hypothetical protein
 hypothetical protein FLJ23282
 early development regulator 1 (homolog of polyomaic 1)
 hypothetical protein PRO2610
 yeast Sec31p homolog
 KIAA0725 protein
 Consensus includes gb:A1246730 /FEA=EST /DB_XREF=gi:3842127 /DB_XREF=est:qk40b01.x1 /CLONE=IMAGE:1871401 /UG=Hs.126901 Homo sapiens mRNA full length i
 diacylglycerol kinase, delta (130kD)
 E1B-55kDa-associated protein 5
 uncharacterized hematopoietic stem/progenitor cells protein MDS030
 RAB30, member RAS oncogene family
 prostaglandin D2 receptor (DP)
 hypothetical protein FLJ10715
 KIAA1735 protein
 thyroid hormone receptor interactor 11
 synaptosomal-associated protein, 91 kDa homolog (mouse)
 cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4)
 RAB32, member RAS oncogene family
 heme-regulated initiation factor 2-alpha kinase
 hypothetical protein FLJ20533
 Consensus includes gb:AL137651.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434O0213 (from clone DKFZp434O0213); partial cds. /FEA=mRNA /GEN=OKFZp434O0213 /F
 ecdonucleotide pyrophosphatase/phosphodiesterase 1
 TP53TG3 protein
 claudin 17
 Consensus includes gb:AK023451.1 /DEF=Homo sapiens cDNA FLJ13389 fts, clone PLACE1001171, moderately similar to MYOTUBULARIN. /FEA=mRNA /DB_XREF=gi:104
 hypothetical protein
 protein x 0001
 major histocompatibility complex, class II, DO alpha
 integrin, beta 4
 letal giant larvae homolog 2 (Drosophila)
 organic cationic transporter-like 4
 Consensus includes gb:BF589529 /FEA=EST /DB_XREF=gi:11681865 /DB_XREF=est:naa0505.x1 /CLONE=IMAGE:3254144 /UG=Hs.106642 ESTs, Weakly similar to hypotr
 sh3-associated polypeptide, 18kD
 pumilo (Drosophila) homolog 2
 KIAA0033 protein
 Consensus includes gb:BF587998 /FEA=EST /DB_XREF=gi:12335213 /DB_XREF=est:602269506F1 /CLONE=IMAGE:4357777 /UG=Hs.21851 Homo sapiens cDNA FLJ1290 BF587998

transducin (beta)-like 1
RCE1 homolog, prenyl protein protease (S. cerevisiae)
secreted protein of unknown function
KIA00943 protein
Consensus includes gb:AL050364.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G052 (from clone DKFZp564G052). /FEA=mRNA /DB_XREF=gi:4914598 /UG=Hs.3027611
deformed epidermal autoregulatory factor 1 (Drosophila)
melanocortin 2 receptor (adrenocorticotrophic hormone)
presenilin 1 (Alzheimer disease 3)
EpNA3
hypothetical protein FLJ10867
pyruvate kinase, liver and RBC
keratin 9 (epidermolytic palmoplantar keratodermia)
Consensus includes gb:AL050376.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586J101 (from clone DKFZp586J101). /FEA=mRNA /DB_XREF=gi:4914609 /UG=Hs.322845 H.
MAP-kinase activating death domain
growth arrest and DNA-damage-inducible, beta
DKFZP568D213 protein
inverdin
DKFZP566C1619 protein
E7A-like factor 1 (eta domain transcription factor)
retinoid acid- and interferon-inducible protein (58kD)
thyroid hormone receptor coactivating protein
sodium channel, voltage-gated, type V, alpha polypeptide (long (electrocardiographic) QT syndrome 3)
G-protein coupled receptor 88
tumor necrosis factor (ligand) superfamily, member 7
SUMO-1-specific protease
melanocortin 5 receptor
hypothetical protein MGC5347
uncharacterized bone marrow protein BM036
PR domain containing 2, with ZNF domain
calcium/calmodulin-dependent protein kinase IV
AT-binding transcription factor 1
KIAA1139 protein
hypothetical protein DKFZp762166
serine/threonine kinase 10
kallitrein 6 (neurosin, zyme)
hepatocellular carcinoma-associated antigen 58
hypothetical protein FLJ13409
epiplatin 1
Consensus includes gb:A1875458 /FEA=EST /DB_XREF=gi:4875938 /DB_XREF=est:wb9911.x.1 /CLONE=IMAGE:2313837 /UG=Hs.206984 Human cosmid CRI-JC2015 at O1
myeloid cell leukemia sequence 1 (BCL2-related)
KIAA1332 protein
tetrahydropeptide repeat domain 4
KIAA0483 protein
G-protein coupled receptor SALPR; somatostatin and angiotensin-like peptide receptor
Consensus includes gb:AL109703.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 25206. /FEA=mRNA /DB_XREF=gi:5689812 /UG=Hs.25314 Hom
BCL2-antagonist of cell death
protocadherin alpha 2
neurexin 2
DKFZP564C186 protein
ortholog of mouse integral membrane glycoprotein LIG-1
phosphate regulating gene with homologies to endopeptidases on the X chromosome (hypophosphatemia, vitamin D resistant rickets)
hypothetical protein FLJ20422
regulator of G-protein signalling 10
KIAA0089 protein
KIAA0700 protein
hypothetical protein MGC5139
protein tyrosine phosphatase, non-receptor type 21
RB-associated KRAB repressor
activin A receptor, type II
H2A histone family, member X
preproenkephalin

AW968555 1.05
NM_005133 1.05
AA156251 1.05
AA53726 1.05
AL050364 1.05
AF088892 1.05
NM_00529 1.05
NM_007319 1.05
NM_005233 1.05
NM_018249 1.05
NM_000298 1.05
NM_000226 1.05
AL050376 1.05
AB002356 1.05
AF078077 1.05
NM_015513 1.05
AF039217 1.05
AL050374 1.05
AL559590 1.05
N47725 1.05
NM_006696 1.05
NM_000335 1.05
NM_022049 1.05
NM_001252 1.05
AK001408 1.05
NM_005913 1.05
AF167079 1.05
AK022348 1.05
NM_018453 1.05
AB61013 1.05
L24959 1.05
NM_006885 1.05
AK025262 1.05
AI341234 1.05
AB015718 1.05
NM_002774 1.05
NM_016436 1.05
NM_024617 1.05
NM_031308 1.05
AI675458 1.05
AI275690 1.05
AI129395 1.05
W22890 1.05
NM_015176 1.05
NM_016566 1.05
AL109703 1.05
U86879 1.05
AF152480 1.05
AK001290 1.05
AB035286 1.05
NM_015658 1.05
AB050468 1.05
U87284 1.05
AI660497 1.05
W19676 1.05
AA135522 1.05
AB014600 1.05
BC004815 1.05
X79510 1.05
NM_021163 1.05
NM_001616 1.05
AA760862 1.05
NM_006228 1.05

Consensus includes gb:221967.1 /DEF=H.sapiens partial cDNA for homologue of mPOU homeobox protein. /FEA=mRNA /PROD=mPOU homeobox protein /DB_XREF=gi:437 Z21967
 abd-interactor 12 (SH3-containing protein) U31089 1.05
 apolipoprotein L AF323540 1.05
 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 5 NM_013252 1.05
 guanylate cyclase 1, soluble, alpha 2 NM_000655 1.05
 phosphorotransferase, muscle U24183 1.05
 hypothetical C2H2 zinc finger protein FLJ22504 AA845577 1.05
 transforming growth factor, alpha NM_003236 1.05
 complement component 4B NM_000592 1.05
 Consensus includes gb:237544 /DEF=H.sapiens gene for phospholipase C beta 3, exon 1 /FEA=mRNA /DB_XREF=gi:609185 /UG=Hs.37121 phospholipase C, beta 3 (phospholipase C, beta 3) Z37544 1.05
 YY1 transcription factor A4748649 1.05
 zinc finger protein 287 AU150728 1.05
 SOX29 AF032454 1.05
 gb:BC000443.1 /DEF=Homo sapiens. Similar to pericentriolar material 1, clone MGC-8458. mRNA, complete cds. /FEA=mRNA /PROD=Similar to pericentriolar material 1 /DB_XREF=gi:10984291 /DB_XREF=est:787g06.x1 /CLONE=IMAGE:3391546 /UG=Hs.130465 ESTs BC000443 1.05
 HBV associated factor BE788439 1.05
 hypothetical protein FLJ20979 NM_024121 1.05
 chromosome 22 open reading frame 3 NM_012285 1.05
 hypothetical protein FLJ13593 NM_024780 1.05
 KIAA0515 protein AK028025 1.05
 methylcrotonyl-Coenzyme A carboxylase 2 (beta) A0439494 1.05
 ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeier-Vogt disease) AF015593 1.05
 BUB1 budding uninhibited by benzimidazoles 1 homolog (yeast) AF043294 1.05
 KIAA0878 protein AK023621 1.05
 Ras association (RAIGDS/AF-6) domain family 1 NM_007182 1.05
 FERM, RhoGEF (ARHGEF) and pleckstrin domain protein 1 (chondrocyte-derived) BF213279 1.05
 fatty acid desaturase 3 NM_021727 1.05
 PAP-1 binding protein BF223045 1.05
 orosomucoid 2 NM_006608 1.05
 histamine receptor H2 NM_022304 1.05
 retinol dehydrogenase 8 (all-trans) NM_015725 1.05
 ADP-ribosyltransferase (NAD+; poly(ADP-ribose) polymerase)-like 2 AK001880 1.05
 caudal type homeo box transcription factor 1 NM_001804 1.05
 general transcription factor IIIC, polypeptide 2 (beta subunit, 110kD) BC000212 1.05
 argininosuccinate synthetase NM_000050 1.05
 KIAA1638 protein NM_025132 1.05
 OXZF2P5668163 protein BC002888 1.05
 Irf upstream element (FUSE) binding protein 3 U89127 1.05
 RAN, member RAS oncogene family BF112006 1.05
 hypothetical protein PP581 NM_025207 1.05
 hypothetical protein FLJ10734 NM_018187 1.05
 susceptibility protein NSG-x NM_016349 1.05
 protease, serine, 2 (trypsin 2) AF009664 1.05
 inositol polyphosphate-1-phosphatase NM_002194 1.05
 hypothetical protein FLJ23558 NM_025095 1.05
 nuclear receptor subfamily 1, group D, member 2 N32659 1.05
 heterogeneous nuclear ribonucleoprotein A1 AL588186 1.04
 chromobox homolog 1 (HP1 beta homolog Drosophila) NM_006807 1.04
 dolichyl-phosphate mannosyltransferase polypeptide 3 NM_018973 1.04
 truncated calcium binding protein NM_016175 1.04
 Consensus includes gb:A172139 /FEA=EST /DB_XREF=gi:4334229 /DB_XREF=est:186a10.x1 /CLONE=IMAGE:2148378 /UG=Hs.9683 DnaJ (Hsp40) homolog, subfamily C, A472139 1.04
 opsin 1 (cone pigments), short-wave-sensitive (color blindness, tritan) NM_001708 1.04
 proteasome (prosome, macropain) 26S subunit, non-ATPase, 1 A1860431 1.04
 TRAF interacting protein NM_005879 1.04
 Consensus includes gb:AK024252.1 /DEF=Homo sapiens cDNA FLJ14190 /fs, clone NT2RP2008534, moderately similar to 5-AMP-ACTIVATED PROTEIN KINASE, CATALYTIC SUBUNIT, isoform 1 AK024252 1.04
 KIAA0644 gene product AV724192 1.04
 hypothetical protein FLJ22968 NM_022842 1.04
 hypothetical protein FLJ12355 NM_024988 1.04
 CASP8 and FADD-like apoptosis regulator U97075 1.04
 T cell receptor delta locus X72501 1.04
 phosphodiesterase 5A, cGMP-specific NM_001083 1.04
 latent transforming growth factor beta binding protein 4 NM_003573 1.04
 shgged-like (fascin homology, sea urchin) (Drosophila) NM_003098 1.04
 Consensus includes gb:BF114815 /FEA=EST /DB_XREF=gi:10984291 /DB_XREF=est:787g06.x1 /CLONE=IMAGE:3391546 /UG=Hs.130465 ESTs BF114815 1.04

KIAA0974 protein	AW300504	1.04
ketohexokinase (fructokinase)	BC006233	1.04
sorting resin 11	NM_013323	1.04
nitric oxide synthase 2A (inducible, hepatocytes)	L24553	1.04
cryptochrome 2 (photolyase-like)	AB014558	1.04
NCX protein	BC006356	1.04
hypothetical protein FLJ20174	NM_017899	1.04
inhibin, beta A (activin A, activin AB alpha polypeptide)	M13436	1.04
KIAA1052 protein	NM_014956	1.04
regulator of G-protein signalling 20	AF074979	1.04
electron-transferring-flavoprotein dehydrogenase	NM_004453	1.04
phospholipase D1, phosphatidylcholine-specific	AL050332	1.04
PHD finger protein 1	NM_002662	1.04
kinectin 1 (kinesin receptor)	Z22551	1.04
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6	NM_016184	1.04
Consensus includes gb:AL109658 /DEF=Human DNA sequence from clone RP4-776F14 on chromosome 20p12.2-13. Contains the 5' end of the FKBP1A gene for FK506-binding protein 1	AL109658	1.04
ubiquitin-activating enzyme E1-like	L13852	1.04
hepatoglobulin	NM_005143	1.04
hypothetical protein FLJ10060	NM_017386	1.04
collagen, type XI, alpha 2	U32168	1.04
phosphoinositide-3-kinase, regulatory subunit 4, p150	BF740111	1.04
lysosomal-associated membrane protein 2	J04183	1.04
gb:NM_014925.1 /DEF=Homo sapiens KIAA1002 protein (KIAA1002). mRNA /FEA=mRNA /GEN=KIAA1002 /PROD=KIAA1002 protein /DB_XREF=gi:7662441 /UG=Hs.10248	NM_014925	1.04
hypothetical protein DKFZp434P2235	AL136860	1.04
hypothetical protein, estradiol-induced	NM_015516	1.04
PRO1914 protein	NM_014106	1.04
Consensus includes gb:U79293.1 /DEF=Human clone 23948 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710274 /UG=Hs.159264 Human clone 23948 mRNA sequence	U79293	1.04
cathepsin E	NM_001910	1.04
aristae-like homeobox 4	NM_021926	1.04
hypothetical protein	AV723665	1.04
ribophorin 1	NM_002950	1.04
collectin sub-family member 10 (C-type lectin)	NM_006438	1.04
pro-melanin-concentrating hormone	NM_002874	1.04
stem cell growth factor; lymphocyte secreted C-type lectin	BC005810	1.04
melastatin 1	NM_002420	1.04
melastatin (EBP72)-like 1	NM_004609	1.04
olfactory receptor, family 10, subfamily J, member 1	NM_012351	1.04
parathyromin	M24398	1.04
hypothetical protein FLJ12150	NM_024736	1.04
karyopherin alpha 5 (importin alpha 5)	NM_002269	1.04
ligase III, DNA, ATP-dependent	NM_002311	1.04
melanoma antigen, family A, 5	AI200443	1.04
soluble carrier family 7 (cationic amino acid transporter, y+ system), member 7	NM_003982	1.04
group-specific component (vitamin D binding protein)	NM_000583	1.04
transcription elongation factor B (SII), polypeptide 1-like	NM_003197	1.04
lipoprotein lipase	NM_000237	1.04
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 3	AI760760	1.04
NAD(P)H dehydrogenase, quinone 2	NM_000904	1.04
kinesin family member 5C	NM_004522	1.04
hypothetical protein FLJ14639	AI854627	1.04
hypothetical protein FLJ13848	NM_024771	1.04
Consensus includes gb:AI821657 /FEA=EST /DB_XREF=gi:5440736 /DB_XREF=est:Zv41809.x5 /CLONE=IMAGE:756184 /UG=Hs.66163 Homo sapiens clone 23636 mRNA	AI821657	1.04
phosphoenolpyruvate carboxykinase 2 (mitochondrial)	NM_004563	1.04
chemokine (C-C motif) receptor 8	NM_005201	1.04
glutathione S-transferase M5	NM_000851	1.04
lectin, galactoside-binding, soluble, 1 (galectin 1)	NM_002305	1.04
spondyloepiphyseal dysplasia, late	NM_014563	1.04
activated leukocyte cell adhesion molecule	BF242805	1.04
Consensus includes gb:AK022303.1 /DEF=Homo sapiens cDNA FLJ12241 /fa. clone MAMMA1001274 /FEA=mRNA /DB_XREF=gi:10433670 /UG=Hs.287503 Homo sapiens	AK022303	1.04
SP110 nuclear body protein	NM_004509	1.04
H326	AK076481	1.04
hypothetical protein FLJ12687	NM_024917	1.04
U2 small nuclear ribonucleoprotein auxiliary factor, small subunit 2	NM_005069	1.04

coagulation factor VIII-associated (intronic transcript) 1.04
 hepatic leukemia factor NM_012151
 nuclear receptor subfamily 4, group A, member 1 1.04
 CASP8 and FADD-like apoptosis regulator NM_003879
 ribonuclease 6 precursor NM_003730
 matrix metalloproteinase 14 (membrane-inserted) 1.04
 hypothetical protein FLJ11753 X83535
 protein kinase D2 NM_024659
 Ellis van Creveld syndrome AF309082
 arginine vasopressin receptor 1A AF216185
 fructose-1,6-bisphosphatase 2 NM_005066
 mitochondrial ribosomal protein L48 NM_003837
 Consensus includes gb:NM55205 /FEA=EST /DB_XREF=gi:1198084 /DB_XREF=est:yy44g05.s1 /CLONE=IMAGE:245624 /UG=Hs.20205 hemoglobin, beta pseudogene 1 NM_016055
 chaperonin containing TCP1, subunit 5 (epsilon) N55205
 LIM domain-containing preferred translocation partner in lipoma AF215798
 PR domain containing 10 AL044018
 5-hydroxytryptamine (serotonin) receptor 2C NM_020228
 zinc finger protein 7 (K0X 4, clone HF.16) NM_000868
 L1 cell adhesion molecule (hydrocephalus, stenosis of aqueduct of Sylvius 1, MASA (mental retardation, aphasia, shuffling gait and adducted thumbs) syndrome, spastic paraparesis) NM_003416
 COL-10 protein AF853981
 killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2 AF23617
 Consensus includes gb:AK026161.1 /DEF=Homo sapiens cDNA: FLJ25508 fts, clone HRC11772. /FEA=mRNA /DB_XREF=gi:10438922 /UG=Hs.8859 Homo sapiens, Similar AK026161
 hypothetical protein FLJ21188 NM_025073
 leptin receptor gene-related protein NM_017526
 Consensus includes gb:AF308291.1 /DEF=Homo sapiens serologically defined breast cancer antigen NY-BR-46 mRNA, partial cds. /FEA=mRNA /PROD=serologically defined AF308291
 TBP-interacting protein NM_018448
 Consensus includes gb:AF308291.1 /DEF=Homo sapiens serologically defined breast cancer antigen NY-BR-46 mRNA, partial cds. /FEA=mRNA /PROD=serologically defined AF308291
 undine phosphorylase NM_003364
 DKFZP566O1648 protein AF201941
 KIAA0788 protein BC001417
 nuclear protein AF273049
 F-box only protein 21 U79257
 Consensus includes gb:AL390736 /DEF=Human DNA sequence from clone RP11-209J19 on chromosome 13 Contains ESTs, STSs and GSSs. Contains the gene for the GWI AL390736 NM_016811
 GPAA1P anchor attachment protein 1 homolog (yeast) BC006883
 zine oculis homeobox homolog 3 (Drosophila) NM_005413
 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1 NM_006162
 v-ort sarcoma virus CT10 oncogene homolog (avian)-like NM_005207
 Consensus includes gb:AA725078 /FEA=EST /DB_XREF=gi:2742785 /DB_XREF=est:a08e12.s1 /CLONE=1342222 /UG=Hs.54567 paired box gene 1 /FL=gb:NM_006192.1 AA725078
 hypothetical protein MGC1203 NM_024296
 immunoglobulin heavy constant mu X17115
 KIAA0682 gene product NM_014562
 wingless-type MMTV integration site family, member 7A O83175
 potassium inwardly-rectifying channel, subfamily K, member 4 NM_016811
 gb:L06101.1 /DEF=Human IG VH-region gene, complete cds. /FEA=mRNA /GEN=IGH@ /PROD=immunoglobulin heavy chain V-region /DB_XREF=gi:185528 /FL=gb:L06101.1 L06101
 transforming growth factor, beta 1 NM_000660
 RYK receptor-like tyrosine kinase X96588
 hypothetical protein MGC5576 NM_024056
 KIAA0237 gene product BC003103
 zinc finger protein 75 (D8C6) W90796
 glycocyphosphatidylinositol specific phospholipase D1 AV699786
 NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 1 (8KD, KFYI) NM_002494
 Consensus includes gb:AK025379.1 /DEF=Homo sapiens cDNA: FLJ21726 fts, clone COLF1088. /FEA=mRNA /DB_XREF=gi:10437882 /UG=Hs.318721 Homo sapiens cDNA: AK025379
 docking protein 1, 82KD (downstream of tyrosine kinase 1) AF180527
 Consensus includes gb:AF0541994.1 /DEF=Homo sapiens clone 23832 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3005709 /UG=Hs.31290 Homo sapiens clone 23832 mRNA AF0541994
 ribosome binding protein 1 homolog 180KD (dog) NM_004587
 hypothetical protein FLJ20535 NM_017868
 myosin IB BF215996
 steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1) BC006373
 hypothetical protein from EUROMIMAGE 1977056 NM_020213
 plasminogen-like AK022897
 reversion-inducing-cytidine-rich protein with kazal motifs NM_004587
 Consensus includes gb:AK025422.1 /DEF=Homo sapiens cDNA: FLJ21769 fts, clone COLF7354. /FEA=mRNA /DB_XREF=gi:10437930 /UG=Hs.306810 Homo sapiens cDNA: AK025422
 phosphatidic acid phosphatase type 2A AF014403
 luronate 2-sulfatase (Hunter syndrome) AF050145

hypothetical protein LOC55565	NM_017530	1.04
cydin 1	BC550368	1.04
hypothetical protein FLJ13224	NM_024799	1.04
8-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 2	NM_008212	1.04
estrogen receptor 2 (ER beta)	AF051428	1.04
creatine kinase, brain	NM_001823	1.04
HIV-1 rev binding protein 2	AI950314	1.04
protein phosphatase 2A, regulatory subunit B' (PR 53)	BC002545	1.04
phospholipase A2, group IIb	AL158172	1.04
hypothetical protein FKSG32	NM_031307	1.04
KIAA0852 protein	NM_014941	1.04
a disintegrin-like and metalloprotease (repolysin type) with thrombospondin type 1 motif, 13	NM_017587	1.04
ribosomal protein S6 kinase, 90kD, polypeptide 2	BC002363	1.04
period homolog 1 (Drosophila)	AF022991	1.04
TGF beta receptor associated protein -1	NM_004257	1.04
KIAA1041 protein	AK027075	1.04
cathepsin W (lymphopain)	NM_001335	1.04
transmembrane protein 1	U61500	1.04
KIAA0414 protein	NM_014007	1.04
hypothetical protein MDS025	NM_021825	1.04
transcription termination factor, RNA polymerase I	NM_007344	1.04
KIAA0202 protein	BC001326	1.04
protein-serine/threonine kinase gene	Z25431	1.04
hypothetical protein FLJ11442	NM_018338	1.04
nuclear receptor subfamily 2, group F, member 1	A195185	1.04
Janus kinase 3 (a protein tyrosine kinase, leukocyte)	U31601	1.04
hypothetical protein	NM_019027	1.04
zinc finger protein 147 (estrogen-responsive finger protein)	NM_005082	1.04
low density lipoprotein receptor (familial hypercholesterolemia)	S70123	1.04
protein phosphatase 1, regulatory (inhibitor) subunit 16B	AB020630	1.04
Rho GTPase activating protein 4	NM_001666	1.04
KIAA0378 protein	Z38645	1.04
similar to ZINC FINGER PROTEIN 85 (ZINC FINGER PROTEIN HPF4) (HTF1) (H. sapiens)	AC006539	1.04
nuclear receptor subfamily 1, group D, member 1	NM_021724	1.04
RAS p21 protein activator (GTPase activating protein) 1	NM_002890	1.04
deoxynucleosyltransferase, terminal	M11722	1.04
protein tyrosine phosphatase, non-receptor type substrate 1	AC004832	1.04
protein kinase C binding protein 1	AB032951	1.04
T-cell leukemia/lymphoma 1B	NM_004918	1.04
thousand and one amino acid protein kinase	NM_004783	1.04
coagulation factor VII-associated (intronic transcript)	AI218431	1.04
Consensus includes gb:AL035287.1 IDEF-H sapiens gene from PAC 747L4 /FEA=mrna /PROD=hypothetical protein /DB_XREF=gi:4200248 /UG=Hs.119254 H sapiens gen	AL035287	1.04
amblyot-sensitive cation channel 2, neuronal	NM_020039	1.04
DKFZP566D193 protein	BE677493	1.04
37 kDa leucine-rich repeat (LRR) protein	NM_005824	1.04
hypothetical protein FLJ20422	NM_017814	1.04
hypothetical protein FLJ12707	NM_022067	1.04
protein kinase, cAMP-dependent, regulatory, type II, alpha	NM_004157	1.04
centaurin, beta 2	D28069	1.04
Consensus includes gb:AA427737 /FEA=EST /DB_XREF=gi:2111576 /DB_XREF=est:zw30g12.s1 /CLONE=IMAGE:770854 /UG=Hs.110820 ESTs	AA427737	1.04
hypothetical protein FLJ20568	NM_017885	1.04
LIM domain kinase 1	D26309	1.04
Consensus includes gb:AI350339 /FEA=EST /DB_XREF=gi:4087545 /DB_XREF=IMAGE:1910362 /UG=Hs.144563 tryptophan hydroxylase (tryptophan	AI350339	1.04
Consensus includes gb:AL567779 /FEA=EST /DB_XREF=gi:12921478 /DB_XREF=est:AL567779 /CLONE=CSODF034YB08 (3 prime) /UG=Hs.84775 Human transposon-like	AL567779	1.04
calmodulin 1 (phosphorylase kinase, delta)	N25325	1.04
protein associated with PRK1	AL136598	1.04
clusterin (complement lytic inhibitor, SP-40,40, sulfated glycoprotein 2, testosterone-repressed prostate message 2, apolipoprotein J)	M25915	1.04
kinasin family member 5B	NM_004521	1.04
tumor necrosis factor (ligand) superfamily, member 5 (hyper-IgM syndrome)	NM_000074	1.04
son of sevenless (Drosophila) homolog 2	AI828605	1.04
acute carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc) transporter), member 3	BC005136	1.04
KIAA1194 protein	NM_015455	1.04
seel-1 suppressor of lin-12-like (C. elegans)	A1927770	1.04

NADH dehydrogenase (ubiquinone) Fe-S protein 1 (75kD) (NADH-coenzyme Q reductase)
 hypothetical protein FLJ14129
 estrogen receptor binding site associated, antigen, 9
 Interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD)
 Consensus includes gb:BF221873 /FEA=EST /DB_XREF=gi:11128850 /DB_XREF=est:7060303.x1 /CLONE=IMAGE:3578429 /UG=Hs.287586 Homo sapiens cDNA FLJ13848 BF221873
 transcriptional regulator interacting with the PHS-bromodomain 2
 erythrocyte membrane protein band 4.1-like 2
 ribosomal protein S14
 receptor-related protein complex 4, sigma 1 subunit
 huntingtin-associated protein 1 (neuroan 1)
 and-associated polypeptide, 18kD
 aldolase B, fructose-bisphosphate
 low density lipoprotein receptor-related protein 6
 hypothetical protein PP1226
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2
 KIAA1074 protein
 transcription factor 2, hepatic; LF-B3; variant hepatic nuclear factor
 penicillin-like 2 (Drosophila)
 e1a variant gene 2
 mitogen-activated protein kinase 14
 chondroitin sulfate proteoglycan 6 (bemarcan)
 inositol 1,4,5-trisphosphate 3-kinase A
 hypothetical protein FLJ12987
 DKFZP564M182 protein
 myosin, light polypeptide 2, regulatory, cardiac, slow
 interferon regulatory factor 5
 thrombospondin 4
 KIAA1072 protein
 cat eye syndrome chromosome region, candidate 1
 hypothetical protein FLJ10521
 gb:L08666.1 /DEF=Homo sapiens porin (por) mRNA, complete cds and truncated cds. /FEA=mrna /GEN=por. por /PROD=porin; porin /DB_XREF=gi:190199 /FL=gb.L08666.1
 U32845
 E74-like factor 4 (e1a domain transcription factor)
 kinesin heavy chain member 2
 v-akt murine thymoma viral oncogene homolog 2
 eukaryotic translation initiation factor 5A2
 prostaglandin E receptor 3 (subtype EP3)
 hypothetical protein FLJ20596
 regulatory factor X, 1 (influences HLA class II expression)
 mucin 4, tracheobronchial
 CLST 11240 protein
 clones Z3887 and Z3775 zinc finger protein
 Consensus includes gb:AL138967 /DEF=Human DNA sequence from clone RP11-149M18 on chromosome 6 Contains ESTs, STSs, GSSs and CpG islands. Contains the natur
 KIAA0392 protein
 RAN binding protein 2
 CDC14 cell division cycle 14 homolog B (S. cerevisiae)
 hypothetical protein DKFZP434G1415
 CASP8 and FADD-like apoptosis regulator
 tripartite motif-containing 31
 Beddus subtilis /REF=X17013 /DEF=B subtilis lys gene for diamino pimelate decarboxylase corresponding to nucleotides 673-1002 of X17013 /LEN=1108 /-5, -M, -3 represent 1
 DKFZP434H132 protein
 macrophage scavenger receptor 1
 Consensus includes gb:BF793446 /FEA=EST /DB_XREF=gi:12098500 /DB_XREF=IMAGE:4347388 /UG=Hs.123122 FSH primary response (LRPF BF793446
 KIAA0268 protein
 granulysin
 protein2008
 Pdu homolog 3 (E. coli)
 paired box gene 4
 acyl-Coenzyme A oxidase 3, pristanoyl
 hypothetical protein FLJ23251
 hypothetical protein similar to preferentially expressed antigen of melanoma
 colony stimulating factor 2 (granulocyte-macrophage)
 Consensus includes gb:AA284903 /FEA=EST /DB_XREF=gi:1927445 /DB_XREF=est:224605.r1 /CLONE=IMAGE:714033 /UG=Hs.306366 Homo sapiens T-cell receptor alpha AA284903
 ligase IV, DNA, ATP-dependent
 NM_005006 1.04
 NM_030895 1.04
 NM_004215 1.04
 BE049439 1.03
 Consensus includes gb:BF221873 /FEA=EST /DB_XREF=est:7060303.x1 /CLONE=IMAGE:3578429 /UG=Hs.287586 Homo sapiens cDNA FLJ13848 BF221873 1.03
 NM_014755 1.03
 BF511685 1.03
 M13934 1.03
 AF155159 1.03
 AF040723 1.03
 BF593650 1.03
 AK026411 1.03
 AF074264 1.03
 AI923458 1.03
 NM_005874 1.03
 AL137351 1.03
 NM_006481 1.03
 U88870 1.03
 AV693985 1.03
 NM_001315 1.03
 AF020043 1.03
 NM_002220 1.03
 NM_025170 1.03
 AK025446 1.03
 AF020768 1.03
 NM_002200 1.03
 NM_003248 1.03
 NM_014906 1.03
 NM_017424 1.03
 BC003073 1.03
 L08666 1.03
 U32845 1.03
 NM_004520 1.03
 AA769075 1.03
 NM_020390 1.03
 D38300 1.03
 AL512707 1.03
 NM_002818 1.03
 NM_004532 1.03
 NM_016438 1.03
 NM_021188 1.03
 W68158 1.03
 AI681120 1.03
 AU145941 1.03
 NM_031292 1.03
 AF041459 1.03
 X61006 1.03
 X17013 1.03
 AW516789 1.03
 AF037351 1.03
 BF793446 1.03
 D87742 1.03
 M85276 1.03
 NM_013395 1.03
 NM_017647 1.03
 NM_006193 1.03
 BF055171 1.03
 NM_024818 1.03
 AL023753 1.03
 M11734 1.03
 Consensus includes gb:AA284903 /FEA=EST /DB_XREF=gi:1927445 /DB_XREF=est:224605.r1 /CLONE=IMAGE:714033 /UG=Hs.306366 Homo sapiens T-cell receptor alpha AA284903 1.03
 NM_002312 1.03

zinc finger protein 36, C3H type-like 1
leukotriene b4 receptor (chemokine receptor-like 1)
nitric oxide synthase 1
Rho guanine exchange factor (GEF) 12
KIAA0097 gene product
KIAA0077 gene product
onc-like 1 (Drosophila)
Consensus includes gb:AL161250 /FEA=EST /DB_XREF=gi:5176917 /DB_XREF=est:w6801.x1 /CLONE=IMAGE:2398489 /UG=Hs.90797 Homo sapiens clone 23620 mRNA : A1761250
myeloid lymphoid or mixed-lineage leukemia (littorax homolog, Drosophila); translocated to 2
cAMP responsive element binding protein 1
neural retina leucine zipper
platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog)
pyruvate dehydrogenase kinase, isoenzyme 3
diacylglycerol kinase, theta (110kD)
myeloid leukemia factor 1
KIAA0276 protein
hypothetical protein MGC13219
hypothetical protein FLJ14153
protein tyrosine phosphatase, receptor type, (polypeptide (PTPRF), interacting protein (liprin), alpha 3
fibroblast growth factor 8 (androgen-induced)
LamC (bacterial lamibiotic synthetase component C)-like 1
KIAA0323 protein
checkpoint with forkhead and ring finger domains
serine/threonine kinase 10
Kell blood group precursor (McLeod phenotype)
RUS guanylate releasing protein 2 (calcium and DAG-regulated)
chemokine (C motif) XC receptor 1
hypothetical protein FLJ20037
zinc finger protein 85 (HPF4, HTF 1)
solite carrier family 16 (monocarboxylic acid transporters), member 1
mitochondrial GTP binding protein
KIAA0542 gene product
FSH primary response (LRPR1 homolog, rat) 1
amelogenin (X chromosome, amelogenesis imperfecta 1)
beta-carotene 15,15'-dioxygenase
nuclear factor of activated T-cells 5, ionically-responsive
tripartite motif-containing 2
perforin 1 (pore forming protein)
solite carrier family 2 (facilitated glucose transporter), member 1
mannose receptor, C type 1
mitochondrial ribosomal protein L18
ATP-binding cassette, sub-family C (CFTR/MRP), member 3
olfactory receptor, family 7, subfamily E, member 38 pseudogene
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 2 (8kD, AGGG)
KIAA0217 protein
hypothetical protein FLJ11585
Consensus includes gb:AF007132.1 /DEF=Homo sapiens clone 23551 mRNA sequence /FEA=mRNA /DB_XREF=gi:2652606 /UG=Hs.184019 Homo sapiens clone 23551 mRNA
KIAA0480 gene product
solite carrier family 22 (organic cation transporter), member 4
Fc fragment of IgG, high affinity 1a, receptor for (CD64)
ALL1-fused gene from chromosome 1q
olfactory receptor, family 7, subfamily A, member 17
DKFZP434F195 protein
zinc finger protein 154 (pH2-92)
Consensus includes gb:AL137145 /DEF=Human DNA sequence from clone RP11.563J2 on chromosome 10 Contains ESTs, STSs, GSSs and a CpG island. Contains a novel AL137145
zinc finger protein 36, C3H type-like 2
Consensus includes gb:AL137474.1 /DEF=Homo sapiens mRNA; cDNA DKFZP434E2423 (from clone DKFZP434E2423) /FEA=mRNA /DB_XREF=gi:6000805 /UG=Hs.24477: AL137474
sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3E
Consensus includes gb:BE045142 /FEA=EST /DB_XREF=gi:8362195 /DB_XREF=est:h28h02.x1 /CLONE=IMAGE:3023283 /UG=Hs.274907 ESTs
fibrogen-like 2
cadherin 6, type 2, X-cadherin (fetal kidney)
platelet-derived growth factor alpha polypeptide
ubiquitin specific protease 9, X chromosome (Drosophila fat facets related)

ephrin-A4	NM_005227	1.03
steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1)	AF113128	1.03
hypothetical protein FLJ10661	NM_018172	1.03
zinc finger protein 287	NM_020853	1.03
CGI-01 protein	NM_015935	1.03
YY1 transcription factor	NM_003403	1.03
hypothetical protein FLJ20241	AI669402	1.03
novel protein	NM_017515	1.03
Consensus includes gb:AK021609.1 /DEF=Homo sapiens cDNA FLJ11547 fs, clone HEMBA1002934, /FEA=mRNA /DB_XREF=gi:10432824 /UG=Hs.306605 Homo sapiens c AK021609	BF081609	1.03
CCAAT/enhancer binding protein (C/EBP), delta	BF081604	1.03
Consensus includes gb:AL060103.1 /DEF=Homo sapiens mRNA; cDNA DKFZ564N2216 (from clone DKFZ564N2216) /FEA=mRNA /DB_XREF=gi:5262528 /UG=Hs.14977	AL060103	1.03
guanine nucleotide binding protein (G protein), alpha activating activity polypeptide, olfactory type	NM_002071	1.03
oncostatin M	BG437034	1.03
deltaX homolog 2 (Drosophila)	AK023924	1.03
macrophage scavenger receptor 1	NM_002445	1.03
brifazrin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome)	X92782	1.03
RAOS homolog (S. pombe)	NM_004584	1.03
hypothetical protein from BCRA2 region	AI609861	1.03
butyrophilin, subfamily 2, member A3	NM_024018	1.03
v-sti sarcoma viral oncogene homolog (avian)	AI588728	1.03
Tax1 (human T-cell leukemia virus type I) binding protein 1	NM_006024	1.03
POU domain, class 2, transcription factor 1	NM_002697	1.03
MUS81 endonuclease	NM_025128	1.03
U2 small nuclear ribonucleoprotein auxiliary factor, small subunit 2	AW089584	1.03
hypothetical protein FLJ12949	NM_023008	1.03
gb:BC004830.1 /DEF=Homo sapiens. Similar to KIAA0798 gene product, clone MGC:10546, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0798 gene product /DB_XREF=gi:10432824	BC004830	1.03
CCAAT/enhancer binding protein (C/EBP), epsilon	NM_001805	1.03
ubiquitin specific protease 20	NM_006876	1.03
tryptase delta 1	NM_012217	1.03
mitogen-activated protein kinase kinase 3	BF971923	1.03
vesicle-associated membrane protein 1 (synaptobrevin 1)	NM_006736	1.03
hypothetical protein FLJ23399	NM_016630	1.03
disvelled, dsh homolog 1 (Drosophila)	NM_022763	1.03
Consensus includes gb:AL133262 /DEF=Human DNA sequence from clone RP1-318D22 on chromosome 6 Contains part of the gene for a CDC5-like protein, the gene for a n	AL133262	1.03
hypothetical protein FLJ10305	NM_018052	1.03
olfactory receptor, family 8, subfamily A, member 1	NM_003696	1.03
MAOS box transcription enhancer factor 2, polypeptide C (myocyte enhancer factor 2C)	NM_002397	1.03
hypothetical protein FLJ21126	NM_025083	1.03
FK506 binding protein 1B (12.6 kD)	AF322070	1.03
radical fringe homolog (Drosophila)	BF940276	1.03
bone morphogenetic protein 6	NM_001718	1.03
cathepsin S	NM_004079	1.03
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3	AF009634	1.03
growth arrest-specific 6	L13720	1.03
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 28	NM_018380	1.03
heat shock transcription factor 1	AI593937	1.03
von Hippel-Lindau binding protein 1	NM_003372	1.03
HSPC034 protein	AV700696	1.03
H2A histone family, member 1	NM_003511	1.03
keratin 10 (epidermolytic hyperkeratosis; keratosis palmaris et plantaris)	M19156	1.03
seroglycan, delta (35kD dystrophin-associated glycoprotein)	NM_000337	1.03
hypothetical protein FLJ10989	AI136679	1.03
discs, large (Drosophila) homolog-associated protein 1	AB000277	1.03
seleconectin W, 1	NM_003009	1.03
breakpoint cluster region	U07000	1.03
hypothetical protein FLJ23329	NM_024958	1.03
PROG14 protein	NM_014131	1.03
kaptein (ectin binding protein)	NM_007059	1.03
hypothetical protein MGC3285	NM_025246	1.03
oxysterol binding protein-like 3	AY008372	1.03
hairy and enhancer of split (Drosophila) homolog 2	NM_019089	1.03

spodysplasia dysplasia, late, pseudogene
 gp:AF130075.1 /DEF=Homo sapiens clone FLB9413 PRO2532 mRNA, complete cds. /FEA=mRNA /PROD=PRO2532 /DB_XREF=gi:11493455 /UG=Hs.306961 Homo sapiens
 hypothetical protein FLJ11781
 KOEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3
 KIAA0254 gene product
 calcitonin receptor-like
 homolog of C. elegans smu-1
 mitogen-activated protein kinase kinase 14
 similar to granzyme B (granzyme 2, cytotoxic T-lymphocyte-associated serine esterase 1) (H. sapiens)
 deoxyhypusine synthase
 KIAA1010 protein
 UDP-Gal-4-epiGalNAc beta 1,4-galactosyltransferase, polypeptide 5
 helicase-mot
 solute carrier family 16 (monocarboxylic acid transporters), member 3
 interferon 8 signal transducer (gp130, oncostatin M receptor)
 protein kinase C binding protein 1
 hemoglobin, alpha 2
 DnaJ (Hsp40) homolog, subfamily B, member 4
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 6
 hypothetical protein FLJ10826
 KIAA0303 protein
 Consensus includes gp:AF339787.1 /DEF=Homo sapiens clone IMAGE:205688, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507325 /UG=Hs.326722 Homo sapiens clone
 pleckstrin homology, Sec7 and coiled-coil domains 4
 hypothetical protein FLJ12270
 Burkitt lymphoma receptor 1, GTP binding protein
 CD28 antigen (Tp44)
 fibronectin II
 Consensus includes gp:AA319460 /FEA=EST /DB_XREF=gi:1971807 /DB_XREF=est:EST21712 /UG=Hs.204166 Human clone 295, 5cm region surrounding hepatocyte nucle
 cortactin SH3 domain-binding protein
 bidata transporter/carrier
 RNA polymerase 1 transcription factor RRN3
 polymerase (DNA directed), beta
 tropomyosin 1 (alpha)
 ubiquitin specific protease
 Consensus includes gp:AW242916 /FEA=EST /DB_XREF=gi:6576686 /DB_XREF=est:xn2703.x1 /CLONE=IMAGE:2694941 /UG=Hs.71988 Homo sapiens mRNA; cDNA DKF
 Rho GTPase activating protein 8
 major histocompatibility complex, class II, DQ alpha 1
 gene near HD on 4p16.3 with homology to hypothetical S. pombe gene
 tubulin-specific chaperone e
 tryptophan rich basic protein
 hypothetical protein
 sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2
 protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform
 Consensus includes gp:AL137534.1 /DEF=Homo sapiens mRNA; cDNA DKFZp34H1419 (from clone DKFZp34H1419); partial cds. /FEA=mRNA /GEN=DKFZp34H1419 /P
 tachykinin receptor 1
 zinc finger protein 124 (HZF-16)
 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)
 lateral giant larvae homolog 1 (Drosophila)
 DKFZP364K247 protein
 mature T-cell proliferation 1
 Consensus includes gp:AI696737 /FEA=EST /DB_XREF=gi:4984637 /DB_XREF=est:wc56102.x1 /CLONE=IMAGE:2322651 /UG=Hs.304757 Homo sapiens similar to human b
 regulator of G-protein signalling 14
 tumor antigen SLP-8p
 hypothetical protein FLJ10251
 sek-1 suppressor of lin-12-like (C. elegans)
 hypothetical protein FLJ11323
 glucosyltransferase 1 (glactoside 2-alpha-L-fucosyltransferase, Bombay phenotype included)
 hypothetical protein MGC10986
 tumor necrosis factor receptor superfamily, member 8
 FYD domain-containing ion transport regulator 3
 RAB27A, member RAS oncogene family
 KIAA0495
 E-1 enzyme

casein kinase 1, gamma 1	AB042563	1.02
transmembrane 4 superfamily member 4	BC001388	1.02
annexin A2	BE908217	1.02
lysine motif-containing 33	AF220137	1.02
guanine nucleotide binding protein (G protein), gamma 5	NM_005274	1.02
aldehyde oxidase 1	NM_001159	1.02
CUG triplet repeat, RNA binding protein 1	NM_006560	1.02
KUAA028 gene product	NM_014724	1.02
adenosine monophosphate deaminase 1 (isoform M)	NM_000036	1.02
T-cell leukemia translocation altered gene	NM_022171	1.02
TONDU	NM_016267	1.02
PDZ domain protein (Drosophila lnaD-like)	NM_005799	1.02
hypothetical protein FLJ11382	NM_021946	1.02
hypothetical protein FLJ10157	NM_018012	1.02
collagen, type V, alpha 1	A1862325	1.02
hypothetical protein FLJ11149	BF340123	1.02
hypothetical protein FLJ20898	NM_024600	1.02
LAP [leucine-rich repeats and PDZ] and no PDZ protein	NM_018214	1.02
axin	AA745954	1.02
v-maf musculosarcoma oncogene homolog F (avian)	AL021977	1.02
hypothetical protein FLJ21432	NM_024551	1.02
transcription factor 8 (represses interleukin 2 expression)	NM_030751	1.02
fetal hypothetical protein	NM_013303	1.02
signal-induced proliferation-associated gene 1	NM_006747	1.02
ATP synthase, H ⁺ transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2	NM_005176	1.02
proteasome (prosome, macropain) inhibitor subunit 1 (PI31)	NM_006814	1.02
leucine-rich repeat protein, neuronal 1	NM_002319	1.02
Consensus includes gb:AA457019 /FEA=EST /DB_XREF=gi:2179739 /DB_XREF=est:aa38a06.s1 /CLONE=IMAGE:815506 /UG=Hs.280645 ESTs. Weakly similar to ALU7_H1	NM_002319	1.02
KUAA0182 protein	D80004	1.02
SRB7 suppressor of RNA polymerase B homolog (yeast)	A1685580	1.02
hypothetical protein FLJ22439	AW027523	1.02
testis derived transcript (3 LIM domains)	BC001451	1.02
hypothetical protein FLJ12122	NM_024979	1.02
Consensus includes gb:AA093357 /FEA=EST /DB_XREF=gi:1645257 /DB_XREF=IMAGE:489600 /UG=Hs.15780 ATP-binding cassette, sub-family A	AA093357	1.02
hypothetical protein FLJ21456	NM_024850	1.02
CD47 antigen (Rb-related antigen, integrin-associated signal transducer)	BG230614	1.02
hypothetical protein CG0003	NM_023037	1.02
Consensus includes gb:A188796 /FEA=EST /DB_XREF=gi:5591960 /DB_XREF=est:wk20b07.x1 /CLONE=IMAGE:2412853 /UG=Hs.243901 Homo sapiens cDNA FLJ20738	A188796	1.02
calcium/calmodulin-dependent protein kinase (CaM kinase) II gamma	AA284757	1.02
KUAA0440 protein	AB007800	1.02
gb:L23519.1 /DEF=Human Ig rearranged gamma-chain, V.DK4-JH4b, complete cds. /FEA=mrna /DB_XREF=gi:388020 /FL=gb:L23519.1	L23519	1.02
hypothetical protein	BG223334	1.02
immunoglobulin kappa variable 1OR2-108	X51887	1.02
protein kinase, cAMP-dependent, catalytic, beta	AA130247	1.02
thrombospondin 1	A1812030	1.02
malate dehydrogenase 2, NAD (mitochondrial)	AL520774	1.02
lipase, endothelial	NM_006033	1.02
melanocortin 3 receptor	NM_019888	1.02
xylosylprotein beta1.4-galactosyltransferase, polypeptide 7 (galactosyltransferase I)	AK022586	1.02
hypothetical protein FLJ23375	NM_024956	1.02
interleukin 1 receptor-like 1	AB012701	1.02
chromosome 1 open reading frame 21	NM_030806	1.02
phosphatidylcholine	AF031463	1.02
solute carrier family 16 (monocarboxylic acid transporters), member 3	AA653175	1.02
HERV-H LTR-associating 3	NM_007071	1.02
killer cell lectin-like receptor subfamily G, member 1	AF081875	1.02
guanylate cyclase 1, soluble, beta 2	NM_004129	1.02
integrin, alpha M (complement component receptor 3, alpha, also known as CD11b (p170), macrophage antigen alpha polypeptide)	J03925	1.02
caspase recruitment domain 4	NM_006092	1.02
ornithine aminotransferase (gyrate atrophy)	NM_000274	1.02
arachidonate lipooxygenase 3	NM_021628	1.02
zinc finger protein 278	A1807017	1.02
hypothetical protein FLJ12438	NM_021833	1.02

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KIAA0258 gene product	NM_014785	1.02
pyruvate dehydrogenase kinase, isoenzyme 3	NM_005391	1.02
NS1-associated protein 1	AF037448	1.02
Consensus includes gb:AK022094.1 /DEF=Homo sapiens cDNA FLJ12032 fls, clone HEMBB1001880. /FEA=mRNA /DB_XREF=gi:10433415 /UG=Hs.151194 Homo sapiens c AK022094	AF354444	1.02
IFP38	AF160249	1.02
KIAA1040 protein	NM_000610	1.02
CD44 antigen (homolog function and Indian blood group system)	BC002752	1.02
PTD015 protein	M12523	1.02
albumin	AA134926	1.02
hypothetical protein FLJ20515	BC002448	1.02
actin binding LIM protein 1	Z83838	1.02
Rho GTPase activating protein 8	NM_017522	1.02
low density lipoprotein receptor-related protein 8, apolipoprotein e receptor	BC000914	1.02
splicing factor, arginine/serine-rich 3	X05610	1.02
collagen, type IV, alpha 2	A051254	1.02
transient receptor potential channel 7	AL136881	1.02
Consensus includes gb:AL359557.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762O1415 (from clone DKFZp762O1415) /FEA=mRNA /DB_XREF=gi:8655612 /UG=Hs.30850	AL136881	1.02
hypothetical protein DKFZp434B044	AL136810	1.02
TBP-interacting protein	NM_018221	1.02
chromosome 2 open reading frame 6	NM_003914	1.02
cyclin A1	NM_016646	1.02
mesenchymal stem cell protein DSCD28	NM_000864	1.02
5-hydroxytryptamine (serotonin) receptor 1D	NM_000505	1.02
coagulation factor XII (Hageman factor)	NM_000016	1.02
acyl-Coenzyme A dehydrogenase, C-4 to C-12 straight chain	NM_002513	1.02
non-metastatic cells 3, protein expressed in	NM_022446	1.02
guanine nucleotide binding protein (G protein), beta polypeptide 1-like	BE543064	1.02
minoxidase, alpha, class 1A, member 2	AK021433	1.02
DKFZP566C243 protein	NM_016099	1.02
HSPC041 protein	NM_004704	1.02
U3 snRNP-associated 55-kDa protein	NM_013317	1.02
lung type-I cell membrane-associated glycoprotein	NM_008760	1.02
uroplakin 2	NM_004452	1.02
estrogen-related receptor beta	AA778936	1.02
Rho guanine nucleotide exchange factor (GEF) 7	A0854869	1.02
E3 ubiquitin ligase SMURF1	AL040538	1.02
protein phosphatase methyltransferase-1	NM_014659	1.02
KIAA0377 gene product	U02520	1.02
collagen, type IV, alpha 3 (Goodpasture antigen)	AL049246	1.02
hypothetical protein FLJ10618	NM_018958	1.02
chromosome 15 open reading frame 2	NM_004087	1.02
chimerin (chimaerin) 2	NM_015850	1.02
mitochondrial ribosomal protein L2	NM_003320	1.02
bubby homolog (mouse)	AW769732	1.02
paired box gene 2	NM_015960	1.02
CGI-32 protein	AW015313	1.02
TATA box binding protein (TBP)-associated factor, RNA polymerase I, C, 110kD	AF001362	1.02
Janus kinase 2 (a protein tyrosine kinase)	AA812224	1.02
KIAA0770 protein	M84073	1.02
asparinyglucosaminidase	NM_002584	1.02
paired box gene 7	NM_001817	1.02
D-amino-acid oxidase	BC001388	1.02
annexin A2	gb:BC002811.1 /DEF=Homo sapiens, Similar to hypothetical protein FLJ20378, clone MGC:3513, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypothetical protein FLJ20378	1.02
calbindin 1, (28kD)	AF049895	1.02
oligodendrocyte lineage transcription factor 2	A0870776	1.02
retinoid acid receptor responder (isazarene induced) 1	A069229	1.02
chromosome 6 open reading frame 34	NM_014320	1.02
inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase gamma	AF091453	1.02
complement component 4A	K02403	1.02
KIAA0914 gene product	AK027138	1.02
uroporphyrinogen decarboxylase	M14016	1.02
DKFZP434B168 protein	AL133049	1.02

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BTG family, member 4	NM_017589	1.02
KIA0461 protein	AB007930	1.02
thorodxin interacting protein	NM_006472	1.02
CGI-115 protein	NM_016052	1.02
v-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian)	AI871354	1.02
pancreatitis-associated protein	NM_002580	1.02
hypothetical protein FLJ10079	NM_017990	1.02
hypothetical protein	BG167522	1.02
core-binding factor, runt domain, alpha subunit 2; translocated to, 1; cyclin D-related	AF018283	1.02
hypothetical protein MGC2603	NM_024037	1.02
immunoglobulin superfamily, member 1	NM_001555	1.02
plexin B1	AV693216	1.02
optic atrophy 1 (autosomal dominant)	BF439570	1.02
ubiquinol-cytochrome c reductase binding protein	M26700	1.02
KIA0578 protein	AB011151	1.02
hypothetical protein FLJ20306	NM_017756	1.02
hypothetical protein FLJ12899	NM_024594	1.02
Consensus includes gb:AL049369.1 /DEF=Homo sapiens mRNA; cDNA DKF7p586D0518 (from clone DKF7p586D0518). /FEA=mRNA /DB_XREF=gi:4500160 /UG=Hs.25072	NM_014788	1.02
tripartite motif-containing 14	NM_001257	1.02
cadherin 13, H-cadherin (heart)	NM_022168	1.02
melanoma differentiation associated protein-5	NM_000723	1.02
calcium channel, voltage-dependent, beta 1 subunit	NM_020231	1.02
xO10 protein	NM_016596	1.02
histone deacetylase 7A	AK024382	1.02
hypothetical protein FLJ14320	NM_004366	1.02
chloride channel 2	NM_001303	1.02
COX10 homolog, cytochrome c oxidase assembly protein, heme A; lamesyltransferase (yeast)	AB046836	1.02
KIAA1816 protein	AB007896	1.02
putative L-type neutral amino acid transporter	AA477655	1.02
Consensus includes gb:AA477655 /FEA=EST /DB_XREF=gi:2206289 /DB_XREF=est:zu37h09.s1 /CLONE=IMAGE:740225 /UG=Hs.181307 H3 histone, family 3A	S82769	1.02
gamma-aminobutyric acid (GABA) A receptor, gamma 3	NM_007372	1.02
RNA helicase-related protein	AF001363	1.02
bridging integrator 1	AV705032	1.02
zinc finger protein 187	NM_018970	1.02
G protein-coupled receptor 85	NM_136967	1.02
Consensus includes gb:AL136967 /DEF=Human DNA sequence from clone RP1.149M18 on chromosome 6 Contains ESTs, STSs, GSSs and CpG islands	NM_005624	1.02
small inducible cytokine subfamily A (Cys-Cys), member 25	NM_014751	1.02
KIAA0429 gene product	L20490	1.02
gamma-glutamyltransferase 1	U08092	1.02
histamine N-methyltransferase	AF070588	1.02
Consensus includes gb:AA456973 /FEA=EST /DB_XREF=gi:2179693 /DB_XREF=est:aa90g11.s1 /CLONE=IMAGE:838628 /UG=Hs.74861 activated RNA polymerase II transcr	AL080191	1.02
hypothetical protein LOC55585	AF007146	1.02
DKF ZP434A062 protein	NM_019009	1.02
Consensus includes gb:AF007146.1 /DEF=Homo sapiens clone 23686 and 23885 mRNA sequences. /FEA=mRNA /DB_XREF=gi:2852624 /UG=Hs.122075 Homo sapiens chr	NM_017842	1.02
Toll-interacting protein	U49396	1.02
hypothetical protein FLJ20489	AF288410	1.02
purinergic receptor P2X, ligand-gated ion channel, 5	NM_024311	1.02
solute carrier family 26, member 6	NM_012417	1.02
Consensus includes gb:AK025352.1 /DEF=Homo sapiens cDNA: FLJ21699 fls, clone COL09829. /FEA=mRNA /DB_XREF=gi:10437851 /UG=Hs.306804 Homo sapiens cDNA: AK025352	NM_005186	1.02
hypothetical protein ET	NM_022578	1.02
retinal degeneration B beta	AF230399	1.01
calpain 1, (mu1) large subunit	NM_004616	1.01
chorionic somatomammotropin hormone-like 1	AU145053	1.01
ADP-ribosylation factor domain protein 1, 64kD	NM_024678	1.01
transmembrane 4 superfamily member 3	BE964689	1.01
formin-binding protein 17	J05084	1.01
hypothetical protein FLJ23441	X97871	1.01
ubiquitin-conjugating enzyme E2L 3	NM_003097	1.01
complement component 6	NM_015997	1.01
erythropoietin receptor	Consensus includes gb:AK021418.1 /DEF=Homo sapiens cDNA FLJ1356 fls, clone HEMBA1000150, highly similar to Homo sapiens putative RNA helicase mRNA. /FEA=mF AK021418	1.01
small nuclear ribonucleoprotein polypeptide N		
CGI-41 protein		

melan-A	U06654	1.01
microspherule protein 1	NM_005337	1.01
Rho-specific guanine nucleotide exchange factor p114	AB011093	1.01
proline-rich Glu (G-carboxyglutamic acid) polypeptide 1	NM_000950	1.01
KIAA0807 protein	AB047005	1.01
potassium channel, subfamily K, member 1 (TWIK-1)	NM_002245	1.01
O-linked mannose beta 1,2-N-acetylglucosaminyltransferase	NM_017739	1.01
cell cycle progression 2 protein	NM_004749	1.01
ring finger protein (C3H2C3 type) 6	AF203342	1.01
hypothetical protein FLJ23040	NM_025174	1.01
hypothetical protein MG5350	AW612374	1.01
Consensus includes gb:AL049278.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564l153 (from clone DKFZp564l153). /FEA=mRNA /DB_XREF=gi:4500033 /UG=Hs.16074 Hon	NM_002090	1.01
GRQ3 oncogene	AW082913	1.01
SFRS protein kinase 1	BC001060	1.01
paired box gene 8	NM_024618	1.01
hypothetical protein FLJ21478	NM_002864	1.01
pregnancy-zone protein	NM_000634	1.01
interleukin 8 receptor, alpha	A040324	1.01
nuclear receptor coactivator 2	A481182	1.01
KIAA0179 protein	AW005776	1.01
DKFZP547E1010 protein	AW974812	1.01
Consensus includes gb:AW974812 /FEA=EST /DB_XREF=gi:6168015 /DB_XREF=est:EST386917 /UG=Hs.291971 ESTs	X17013	1.01
Bacillus subtilis /REF=X17013 /DEF=B subtilis lys gene for diaminopimelate decarboxylase corresponding to nucleotides 1008-1263 of X17013 /LEN=1108 (-S, -M, -3 represent	D87486	1.01
KIAA0278 protein	AW242297	1.01
microtubule-associated protein 7	NM_014178	1.01
HSPC156 protein	AF142482	1.01
TEA domain family member 3	AA527515	1.01
Cluster incl. AA527515:ng11h08.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:937403 /clone_end=3 /gb=AA527515 /gi=2289584 /ug=Hs.238570 /len=587	AF042438	1.01
smoothelin	U85658	1.01
transcription factor AP-2 gamma (activating enhancer binding protein 2 gamma)	D63877	1.01
KIAA0157 protein	NM_007321	1.01
RAN binding protein 3	AW025529	1.01
serologically defined breast cancer antigen NY-BR-20	NM_022105	1.01
death associated transcription factor 1	NM_015833	1.01
adenosine deaminase, RNA-specific, B1 (homolog of rat REDI1)	U50277	1.01
Consensus includes gb:U50277.1 /DEF=Human breast cancer suppressor element (shmael Upper CP1 mRNA, partial cds. /FEA=mRNA /PROD=suppressor element (shmael	NM_006319	1.01
COP-diacylglycerol-inositol 3-phosphatidylinositol synthase (phosphatidylinositol synthase)	A1148659	1.01
Consensus includes gb:A1148659 /FEA=EST /DB_XREF=gi:3677128 /DB_XREF=est:qd69c01.x1 /CLONE=IMAGE:1714848 /UG=Hs.296328 ESTs	NM_018036	1.01
hypothetical protein FLJ10242	NM_000531	1.01
ornithine carbamoyltransferase	BC000941	1.01
SELENOPHOSPHATE SYNTHETASE ; Human selenium donor protein	NM_020234	1.01
x ODR protein	M87313	1.01
dystrophia myotonica-protein kinase	BC003666	1.01
zinc finger protein 24 (K0X 17)	AL516159	1.01
ubiquitin-conjugating enzyme E2E 1 (UBC4/5 homolog, yeast)	BF340228	1.01
insulin-like growth factor binding protein 3	U50383	1.01
retinoic acid induced 15	NM_002987	1.01
small inducible cytokine subfamily A (Cys-Cys), member 17	NM_018158	1.01
solute carrier family 4 (anion exchanger), member 1, adaptor protein	NM_018107	1.01
hypothetical protein FLJ10482	U53470	1.01
inositol polyphosphate-5-phosphatase, 145KD	NM_011395	1.01
Consensus includes gb:BE531136 /FEA=EST /DB_XREF=gi:9759781 /DB_XREF=est:601278315F1 /CLONE=IMAGE:3610539 /UG=Hs.8982 ESTs, Highly similar to KIAA1136	NM_014662	1.01
KIAA0645 gene product	AF119418	1.01
sialyltransferase 9 (CMP-NeuAc: lactosylceramide alpha-2,3-sialyltransferase; GM3 synthase)	AU146781	1.01
Consensus includes gb:AU146781 /FEA=EST /DB_XREF=gi:11008312 /DB_XREF=est:AU146781 /CLONE=HEMBB1001531 /UG=Hs.287474 Homo sapiens cDNA FLJ12000	BE56378	1.01
lipidoxin	BC008005	1.01
gp:AF249277.1 /DEF=Homo sapiens cervical cancer suppressor-1 mRNA, complete cds. /FEA=mRNA /PROD=cervical cancer suppressor-1 /DB_XREF=gi:10442005 /UG=Hs.	BF515124	1.01
uncharacterized hematopoietic stem/progenitor cells protein MDS032	BC000373	1.01
KIAA0828 protein	NM_006333	1.01
amyloid beta (A4) precursor-like protein 2	AK021457	1.01
nuclear DNA-binding protein	NM_005198	1.01
Consensus includes gb:AK021457.1 /DEF=Homo sapiens cDNA FLJ11395 (fs, clone HEMBA1000594. /FEA=mRNA /DB_XREF=gi:10432649 /UG=Hs.306595 Homo sapiens c		
choline kinase-like		

pre-B-cell leukemia transcription factor 3	NM_006195
bridging integrator 1	AF043899
KIAA0239 protein	NM_015288
choline acetyltransferase	NM_020985
phosphatidylinositol-4-phosphate 5-kinase, type I, alpha	U76577
Fanconi anemia, complementation group A	BC048069
minor histocompatibility antigen HB-1	NM_021182
sedc 82 kDa protein mRNA	NM_014597
hypothetical protein FLJ10782	NM_018216
G protein-coupled receptor 6	U18549
membrane component, chromosome 17, surface marker 2 (ovarian carcinoma antigen CA125)	NM_005899
hypothetical protein FLJ13055	NM_022737
cydin-dependent kinase (CDC2-like) 10	NM_003674
ecotropic viral integration site 5	AF008915
proteasome (prosome, macropain) subunit, alpha type, 2	AL523904
cathepsin B	NM_001908
topoisomerase (DNA) III alpha	NM_004618
splicing factor, arginine/serine-rich 2, interacting protein	AW084759
thromboxane A synthase 1 (platelet, cytochrome P450, subfamily V)	NM_030984
Sec23-interacting protein p125	AI322982
fumarylacetoacetate hydrolase (fumarylacetoacetase)	NM_000137
SP110 nuclear body protein	AF280094
hypothetical protein FLJ22353	NM_024587
glucuronidase, beta	NM_000181
mammian-binding lectin serine protease 1 (C4/C2 activating component of Ra-reactive factor)	BC000587
E1A binding protein p300	AI459462
DKFZP964A122 protein	NZ4643
killer cell lectin-like receptor subfamily C, member 3	NM_002261
gb:NM_016473.1 /DEF=Homo sapiens hypothetical protein (LOC51243), mRNA. /FEA=CDS /GEN=LOC51243 /PROD=hypothetical protein /DB_XREF=gi:7706014 /UG=Hs.26	NM_016473
KIAA0700 protein	BC001205
hypothetical protein FKSG63	NM_031221
HSPCO34 protein	AL031427
KIAA0877 protein	AK002054
chromosome X open reading frame 12	NM_003492
gb:NM_001119.2 /DEF=Homo sapiens adducin 1 (alpha), transcript variant 1, mRNA. /FEA=mrna /GEN=adducin 1 (alpha) isoform a /DB_XREF=gi:77060119	NM_001119
DHHC1 protein	NM_016598
PCF11p homolog	AB020631
praja 1	NM_022368
enigma (LIM domain protein)	AI825846
high-mobility group 20B	NZ5429
pellet pan homolog (Drosophila)	BC000535
KIAA0700 protein	AB014600
enhancer of zeste homolog 2 (Drosophila)	NM_004456
hypothetical protein FLJ11280	AL561943
hypothetical protein FLJ11712	NM_024570
putative N-acetyltransferase Camello 2	NM_016347
hypothetical protein FLJ10213	NM_018029
nucleolar cysteine-rich protein	BC004421
glutamate-ammonia ligase (glutamine synthase)	U08626
nucleophosmin (nucleolar phosphoprotein B23, numatrin)	AA191576
aldolase B, fructose-bisphosphate	AF652403
Consensus includes gb:AF098968.1 /DEF=Homo sapiens familial Mediterranean fever focus region, mRNA sequence. /FEA=mrna /DB_XREF=gi:4323318 /UG=Hs.131111 H.AF098968	N39536
p45 protein	gb:Z25422.1 /DEF=H.sapiens protein-serine/threonine kinase /DB_XREF=gi:405772 /FL=gb.Z25422.Z25422
peroxiredoxin 2	L19185
homogentisate 1,2-dioxygenase (homogentisate oxidase)	AI478172
Consensus includes gb:AL109698.1 /DEF=Homo sapiens mRNA (full length insert cDNA clone EUROMAGE 21920). /FEA=mrna /DB_XREF=gi:5689771 /UG=Hs.185701	Horr AL109698
peroxisomal fatty acylated protein	BC000496
slit homolog 3 (Drosophila)	NM_003062
glutamate receptor, ionotropic, AMPA 3	NM_007325
keratin 18	AL022068
nicotinamide N-methyltransferase	NM_006169
solute carrier family 8 (neurotransmitter transporter, tauroine), member 6	NM_003043

KIAA0487 protein
 Consensus includes gb:AA741028 FEA=EST /DB_XREF=gi:2779620 /DB_XREF=est:ny99c07.s1 /CLONE=IMAGE:1286412 /UG=Hs.256155 ESTs. Moderately similar to ALU. AA741028
 hypothetical protein FLJ20262
 tumor necrosis factor, alpha-induced protein 1 (endothelial)
 gqg1 autoantigen, golgin subfamily a, 1
 kinesin family member 3C
 hypothetical protein PRO2984
 transcription elongation factor 8 (SUI), polypeptide 3 (110kD, elongin A)
 Consensus includes gb:AK023081.1 /DEF=Homo sapiens cDNA FLJ13019 fis. clone NT2RP3000738, highly similar to Human mRNA for KIAA0140 gene. /FEA=mRNA /DB_XJAK023081
 surfactant, pulmonary-associated protein C
 schaefer-scute complex-like 1 (Drosophila)
 cytochrome P450, subfamily IIIA (naphthalene oxidase), polypeptide 4
 Consensus includes gb:AA845825 FEA=EST /DB_XREF=gi:2931965 /DB_XREF=est:ak85f12.s1 /CLONE=IMAGE:1414703 /UG=Hs.282721 ESTs. Weakly similar to ALU7_HJAA845825
 solute carrier family 15 (oligopeptide transporter), member 1
 Consensus includes gb:AL134612 /FEA=EST /DB_XREF=gi:6602799 /DB_XREF=est:DKFZp547E086.s1 /CLONE=DKFZp547E086 /UG=Hs.167038 Homo sapiens clone 237 AL134612
 SMA5
 protein inhibitor of activated STAT3
 lifeguard
 hypothetical protein FLJ11181
 periaxin
 cholinergic receptor, nicotinic, epsilon polypeptide
 neuregulin 1
 cytokine-like nuclear factor n-pac
 G protein-coupled receptor, family C, group 5, member B
 hypothetical protein R33729.1
 bone morphogenetic protein.1
 hypothetical protein FLJ20506
 KIAA0411 gene product
 p47 protein
 procollagen-proline, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide I
 myosin, heavy polypeptide 11, smooth muscle
 Eap-related gene member 2
 MLL septin-like fusion
 ret finger protein-like 3 antisense
 spectrin, beta, non-erythrocytic 1
 mitogen-activated protein kinase 8 interacting protein 3
 chromosome 21 open reading frame 2
 hypothetical protein FLJ23282
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide, Y chromosome
 surfactant, pulmonary-associated protein C
 small inducible cytokine subfamily E, member 1 (endothelial monocyte-activating)
 epilipoprotein L, 6
 protein tyrosine phosphatase, receptor type, C
 phospholipase A2, group X
 ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease)
 BH-proteadherin (brain-heart)
 mesenchymal stem cell protein DSC43
 p21(CDKN1A)-activated kinase 4
 keratin 16 (focal non-epidermolytic palmoplantar keratoderma)
 anti-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phospholipase A2)
 hypothetical protein FLJ23322
 caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)
 NADPH oxidase 4
 KIAA0295 protein
 keratin 10 (epidermolytic hyperkeratosis; keratosis palmaris et plantaris)
 integrin, alpha 2b (platelet glycoprotein IIb of fibrin complex, antigen CD41B)
 hypothetical protein FLJ20083
 centrosomal protein 1
 progesterone-associated endometrial protein (placental protein 14, pregnancy-associated endometrial alpha-2-globulin, alpha uterine protein)
 myeloid/lymphoid or mixed-lineage leukemia 2
 sigma receptor (SR31747 binding protein 1)
 KIAA1233 protein
 calcium channel, voltage-dependent, beta 3 subunit

mitogen-activated protein kinase 11	U53442	1.01
HSPC157 protein	NM_014179	1.01
hypothetical protein FLJ20598	NM_017895	1.01
hypothetical protein from clones 23549 and 23762	NM_021226	1.01
sperm autoantigenic protein 17	NM_017425	1.01
chemokine (C-C motif) receptor 1	A1421071	1.01
suc1-associated neurotrophic factor target 2 (FGFR signalling adaptor)	NM_008653	1.01
hypothetical protein MG24171	NM_024307	1.01
angiotensin receptor 1	NM_000685	1.01
homeo box B7	AW102783	1.01
hypothetical protein FLJ21156	NM_024602	1.01
transient receptor potential channel 2	A1759824	1.01
ubiquitin-cytochrome c reductase binding protein	NM_006294	1.01
ATPase, aminophospholipid transporter-like, Class I, type 8A, member 2	AU148927	1.01
KIAA0864 protein	AK025604	1.01
potassium inwardly-rectifying channel, subfamily J, member 4	U24056	1.01
titin immunoglobulin domain protein (myotilin)	NM_008790	1.01
G protein-coupled receptor 17	Z94154	1.01
cytoplasmic FMRP interacting protein 2	AL161999	1.01
KIAA1922 protein	AK026910	1.01
caldesmon 1	A185060	1.01
ketohexokinase (fructokinase)	NM_000221	1.01
KIAA0668 protein	AK021690	1.01
retinal degeneration, slow (retinitis pigmentosa 7)	NM_000322	1.01
enigma (LIM domain protein)	NM_005451	1.01
Tcd37 homolog	AF123538	1.01
protein kinase C-like 2	AF118089	1.01
Consensus includes gb:U80770.1 /DEF=Human EST clone 251800 mariner transposon Hsmar1 sequence. /FEA=mrna /DB_XREF=gi:2921608 /UG=Hs.42192 Human EST cl U80770	U80770	1.01
hypothetical protein 689	BC000819	1.01
KIAA0710 gene product	NM_014871	1.01
Ellis van Creveld syndrome	NM_014556	1.01
heat shock 105kD	BC403660	1.01
signal recognition particle 14kD (homologous Alu RNA binding protein)	NM_003134	1.01
vesicle-associated membrane protein 5 (myobrevin)	NM_006634	1.01
mitochondrial ribosomal protein S4	AL138913	1.01
inner centromere protein antigens (135kD, 155kD)	NM_020238	1.01
polymerase (DNA directed), lambda	NM_013274	1.01
mannosyl (alpha-1,6)-glycoprotein beta-1,6-N-acetyl-glucosaminyltransferase	NM_002410	1.01
peptidase D	NM_000285	1.01
hypothetical protein P15-2	AF201942	1.01
DiGeorge syndrome critical region gene 2	H42875	1.01
Consensus includes gb:A1273812 /FEA=EST /DB_XREF=gi:3896080 /DB_XREF=astqu08107.x1 /CLONE=IMAGE:1964197 /UG=Hs.169252 Human L-myc protein gene, comp A1273812	A1273812	1.01
peptidylglycine alpha-amidating monooxygenase	A022862	1.01
Consensus includes gb:AV728268 /FEA=EST /DB_XREF=gi:10837689 /DB_XREF=eat:AV728268 /CLONE=HTCACH05 /UG=Hs.82845 Homo sapiens cDNA: FLJ21930 fls, ct AV728268	AV728268	1.01
Consensus includes gb:AK025206.1 /DEF=Homo sapiens cDNA: FLJ21553 fls, clone COL06328. /FEA=mrna /DB_XREF=gi:10437671 /UG=Hs.308786 Homo sapiens cDNA: AK025206	AK025206	1.01
collagen, type IV, alpha 3 (Goodpasture antigen)	U02520	1.01
selectin P ligand	U02287	1.01
hypothetical protein FLJ21817 similar to Rho1p2	NM_022448	1.01
heat shock 27kD protein 3	NM_006308	1.01
HSPC047 protein	NM_014147	1.01
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	AL117663	1.01
KIAA0779 protein	AB018322	1.01
feline sarcoma oncogene	NM_002005	1.01
G-2 and S-phase expressed 1	BF973178	1.01
leucine carboxyl methyltransferase	SC004214	1.01
POU domain, class 6, transcription factor 1	L14482	1.01
hypothetical protein FLJ12577	NM_030925	1.01
STAT induced STAT inhibitor 3	NM_003955	1.01
ribosomal protein L39	BC001019	1.00
ubiquitin 3	NM_017481	1.00
hypothetical protein FLJ13993	NM_024695	1.00
thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)	X55005	1.00
hypothetical protein	A1800419	1.00

olfactory receptor, family 1, subfamily E, member 1
 cadherin 2, type 1, N-cadherin (neuronal)
 calcium channel, voltage-dependent, alpha 1F subunit
 RAB27B, member RAS oncogene family
 adaptor-related protein complex 3, sigma 2 subunit
 leukocyte membrane antigen
 retinoic acid receptor responder (tazarotene induced) 1
 thyrotropin-releasing hormone
 protocadherin gamma subfamily A, 11
 ras-related C3 botulinum toxin substrate 2 (rho family, small GTP binding protein Rac2)
 SKI-like
 adaptor-related protein complex 2, alpha 2 subunit
 hypothetical protein FLJ22609
 UV radiation resistance associated gene
 cannabinoid receptor 1 (brain)
 KIAA0395 protein
 peroxisomal biogenesis factor 7
 luciferase transferase 2 (secretor status included)
 Ewing sarcoma breakpoint region 1
 Consensus includes gb:AJ275397 /DEF=Homo sapiens partial (GVH1 gene for immunoglobulin heavy chain V region, case 1, cell Mo V 94 /FEA=CDS /DB_XREF=gi:7573056 /AJ275397
 G protein-coupled receptor
 gb:NM_003871.1 /DEF=Homo sapiens CDC14 (cell division cycle 14, S. cerevisiae) homolog B (CDC14B), mRNA, /FEA=mRNA /GEN=CDC14B /PROD=S, cerevisiae CDC14
 synovial sarcoma, X breakpoint 4
 KIAA1023 protein
 unoplectin 1A
 hypothetical protein FLJ10925
 RAB3A interacting protein (rab3n3)-like 1
 caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)
 polymerase (DNA directed), theta
 hypothetical protein FLJ11137
 protocadherin alpha 5
 N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1
 aminopeptidase
 deleted in cancer 1; RNA helicase HDB/DICE1
 ribosomal protein L23a-like 1
 T-box, brain, 1
 fetal Alzheimer antigen
 transducin-like enhancer of split 1 (E(sp1) homolog, Drosophila)
 hypothetical protein FLJ10248
 methionyl aminopeptidase 2
 actin cross-linking factor
 bromodomain adjacent to zinc finger domain, 1B
 inositol 1,4,5-trisphosphate receptor, type 2
 CCG triplet repeat binding protein 1
 natural killer cell transcript 4
 Consensus includes gb:AA129909 /FEA=EST /DB_XREF=gi:1689667 /DB_XREF=est:zn8808.r1 /CLONE=IMAGE:565091 /UG=Hs.145357 ESTs, Moderately similar to ALU7_
 hypothetical protein FLJ20127
 KIAA0470 gene product
 protocadherin 11
 Consensus includes gb:AT34111 /FEA=EST /DB_XREF=gi:5055224 /DB_XREF=est:ncd8601.y5 /CLONE=IMAGE:771984 /UG=Hs.115670 ESTs
 gb:NM_024304.1 /DEF=Homo sapiens hypothetical protein MGC4859 similar to HSPA8 (MGC4859), mRNA, /FEA=mRNA /GEN=MGC4859 /PROD=hypothetical protein MGC
 Consensus includes gb:AL049307.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564A216 (from done DKFZp564A216) /FEA=mRNA /DB_XREF=gi:4500076 /UG=Hs.274505 t-AL049307
 synaptosomal-associated protein, 25kD
 Consensus includes gb:AY963138 /FEA=EST /DB_XREF=gi:8152974 /DB_XREF=est:EST375211 /UG=Hs.302040 Homo sapiens DNA sequence from PAC 434O14 on chr9
 Consensus includes gb:AV721967 /FEA=EST /DB_XREF=est:AV721967 /CLONE=HTBAPB10 /UG=Hs.184779 Homo sapiens mRNA; cDNA DKFZp5
 Bacillus subtilis /REF=X04603 /DEF=B subtilis thc. thcB genes corresponding to nucleotides 1689-2151 of X04603 /LEN=2073 (-5, -M, -3 represent transcript regions 5 prime, X04603
 homeo box A4
 hypothetical protein FLJ11110
 steroid carrier protein 2
 Consensus includes gb:AW268081 /FEA=EST /DB_XREF=gi:6655111 /DB_XREF=est:xp92e03.x1 /CLONE=IMAGE:2767612 /UG=Hs.193408 ESTs, Weakly similar to ALUC_
 bridging integrator 1
 dual-specificity lysine-(Y)-phosphorylation regulated kinase 1A
 D86550

DnaJ (Hsp40) homolog, subfamily B, member 12
 adenylyate kinase 3
 fragile X mental retardation 1
 hypothetical protein FLJ20583
 carboxypeptidase A2 (pancreatic)
 hypothetical protein PRQ2015
 VRK3 for vacuolar related kinase 3
 proliferation-associated ZGA, 38kD
 Consensus includes gb:M20619 /DEF=Human DNA with a hepatitis B virus surface antigen (HBsAg) gene (complete cds).insertion /FEA=CDS /DB_XREF=gi:184331 /UG=Hs.1 M20619
 calcium binding protein P22
 carbohydrate (N-acetylglucosamine-6-O) sulfotransferase 2
 Consensus includes gb:AB041269.1 /DEF=Homo sapiens mRNA for keratin 19, partial cds, isolate K19-141. /FEA=mRNA, (GEN=keratin 19 /PROD=keratin 19 /DB_XREF=gi:77:AB041269
 latexin protein
 T cell receptor delta locus
 KIAA0116 protein
 HSPCO34 protein
 pM5 protein
 Consensus includes gb:BE845861 /FEA=EST /DB_XREF=gi:970172 /DB_XREF=est:7677d05.x1 /CLONE=IMAGE:3288469 /UG=Hs.24064 Homo sapiens mRNA: cDNA DKF B6645861
 interleukin 1 receptor, type II
 peroxisomal biogenesis factor 7
 HIC1-related gene on chromosome 22
 hypothetical protein FLJ21841
 nuclear receptor subfamily 2, group F, member 2
 collagen, type VI, alpha 1
 sorbitol dehydrogenase
 FK506 binding protein 18 (12.8 kD)
 ankylosis, progressive homolog (mouse)
 CGI-98 protein
 potassium voltage-gated channel, shaker-related subfamily, member 4
 chromosome 21 open reading frame 2
 DNA-binding transcriptional activator
 cytochrome P450, subfamily X2A (steroid 21-hydroxylase, congenital adrenal hyperplasia), polypeptide 2
 Consensus includes gb:AK025191.1 /DEF=Homo sapiens cDNA: FLJ21538 fts, clone COL06151. /FEA=mRNA /DB_XREF=gi:10437854 /UG=Hs.306783 Homo sapiens cDNA: AK025191
 sperm associated antigen 8
 sulfotransferase family 4A, member 1
 potassium inwardly-rectifying channel, subfamily J, member 6
 dolichyl-phosphate mannosyltransferase polypeptide 2, regulatory subunit
 eukaryotic translation initiation factor 4E-like 3
 heterogeneous nuclear ribonucleoprotein A1
 dihydropyrimidinase
 Consensus includes gb:AL139349 /DEF=Human DNA sequence from clone RP11-281P9 on chromosome 20. Contains ESTs, STSs, GSSs and a CpG island. Contains the 3 p: AL139349
 ATPase, Ca++ transporting, type 2C, member 1
 NP0009 protein
 dopamine beta-hydroxylase (dopamine beta-monoxygenase)
 transition protein 2 (during histone to protamine replacement)
 similar to rat myomegalin
 cyclin T1
 hypothetical protein FLJ11040
 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2
 interleukin-1 receptor-associated kinase M
 gamma-tubulin complex protein 2
 Cluster Ind. A1186666: q62a10.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:1745466 /gi=3737304 /ug=Hs.21497 /len=512
 PTPRF Interacting protein, binding protein 2 (lpirin beta 2)
 hypothetical protein FLJ22813
 NCK-associated protein 1
 solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2

ATP synthase, H⁺ transporting, mitochondrial F1 complex, gamma polypeptide 1
 KIAA0586 gene product
 CGI-58 protein
 fragile X mental retardation 2
 protein kinase C, delta
 polo (Drosophila)-like kinase
 glycerol kinase
 hypothetical protein FLJ13782
 peroxisomal biogenesis factor 16
 KIAA0268 protein
 transmembrane 6 superfamily member 1
 protein phosphatase 2, regulatory subunit B (B56), epsilon isoform
 gp:NM_002913.1 /DEF=Homo sapiens replication factor C (activator 1) 1 (145KD) (RFC1), mRNA, /FEA=mRNA /GEN=RFC1 /PROD=replication factor C (activator 1) 1 (145KD) (RFC1)
 KIAA0978 protein
 chromosome 11 open reading frame 5
 signal transducer and activator of transcription 1, 91KD
 RNA polymerase I subunit
 RAE1 RNA export 1 homolog (S. pombe)
 developmentally regulated GTP binding protein 2
 hypothetical protein FLJ20373
 lysyl oxidase-like 2
 NRAS-related gene
 interleukin enhancer binding factor 1
 hypothetical protein FLJ12816
 matrix metalloproteinase 15 (membrane-inserted)
 general transcription factor IIIA
 hypothetical protein MGC5378
 RAD23 homolog 6 (S. cerevisiae)
 aldehyde dehydrogenase 3 family, member B2
 splicing factor 3b, subunit 1, 155KD
 adenyl cyclase-associated protein 2
 Consensus includes gb:AL034550 /DEF=Human DNA sequence from clone RPS-1184F4 on chromosome 20q11.1-11.23. Contains the 3' end of gene KIAA0978, two genes for AL034550
 gb:U95006.1 /DEF=Human D9 splice variant A mRNA, complete cds, /FEA=mRNA /PROD=D9 splice variant A /DB_XREF=gi:2071992 /UG=Hs.37616 Human D9 splice variant
 KIAA0062 protein
 transforming growth factor, beta receptor II (70-80KD)
 raslike protein
 HGC8.1.1 protein
 hypothetical protein
 KIAA1332 protein
 LDL induced EC protein
 NS1-binding protein
 hypothetical protein RG083M05.2
 hypothetical protein MGC3121
 Consensus includes gb:BF001666 /FEA=EST /DB_XREF=gi:10701941 /DB_XREF=est:7g81d12.x1 /CLONE=IMAGE:3313847 /UG=Hs.12460 Homo sapiens clone 23870 mRN
 dual specificity phosphatase 6
 chromosome 21 open reading frame 66
 Consensus includes gb:A928508 /FEA=EST /DB_XREF=gi:3076797 /DB_XREF=estom17g03.s1 /CLONE=IMAGE:1541332 /UG=Hs.10762 ESTs
 DKFZP564A2418 protein
 Ras-GTPase-activating protein SH3-domain-binding protein
 solute carrier family 12 (potassium/chloride transporters), member 7
 hypothetical protein FLJ11132
 DR1-associated protein 1 (negative cofactor 2 alpha)
 H2B histone family, member A
 azurophilin
 hypothetical protein FL10818
 karyopherin (importin) beta 1
 SA (rat hypertension-associated) homolog
 cargo selection protein (mannose 6 phosphate receptor binding protein)
 hypothetical protein MGC14799
 nipasap homolog 1 (C. elegans)
 erythrocyte membrane protein band 4.1-like 3
 inhibitor of growth family, member 1
 golgi SNAP receptor complex member 2

BC000931 1,24
 AB011158 1,24
 NM_016408 1,24
 AJ001550 1,24
 NM_008254 1,24
 NM_005030 1,24
 NM_000167 1,24
 NM_024815 1,24
 AA523441 1,24
 AB006624 1,24
 NM_023003 1,24
 NM_008246 1,24
 NM_002813 1,24
 AL117518 1,24
 NM_014205 1,24
 M87835 1,24
 AF008442 1,24
 NM_003610 1,24
 BF223206 1,24
 NM_017792 1,24
 BE251211 1,24
 NM_007156 1,23
 NM_004514 1,23
 NM_022060 1,23
 NM_002428 1,23
 BE542815 1,23
 BC000927 1,23
 NM_002874 1,23
 AA071510 1,23
 AW003030 1,23
 N90755 1,23
 Consensus includes the 3' end of gene KIAA0978, two genes for AL034550
 U95006 1,23
 C31887 1,23
 D50683 1,23
 BF978689 1,23
 AB016900 1,23
 NM_023070 1,23
 AA636114 1,23
 NM_016202 1,23
 AB020657 1,23
 AC000064 1,23
 NM_024031 1,23
 BC005047 1,23
 NM_013329 1,23
 AA928508 1,23
 AK002064 1,23
 NM_005754 1,23
 NM_006598 1,23
 NM_018335 1,23
 NM_006442 1,23
 BE271470 1,23
 BC003404 1,23
 AB27844 1,23
 BG249565 1,23
 AC004381 1,23
 NM_005817 1,23
 BC005995 1,23
 NM_003634 1,23
 NM_012307 1,23
 NM_005537 1,23
 AW148492 1,23

titin
 nuclear receptor subfamily 3, group C, member 1
 hypothetical protein DKFZp434B195
 hypothetical protein FLJ20030
 G protein-coupled receptor 8
 Consensus includes gb:BF448315 /FEA=EST /DB_XREF=gi:11513897 /DB_XREF=est:mad18a03.x1 /CLONE=IMAGE:3365668 /UG=Hs.152939 Homo sapiens clone 24630 m /BF448315
 KIAA0355 gene product
 ankyrin 3, node of Ranvier (ankyrin G)
 nuclear transcription factor, X-box binding 1
 prolactin regulatory element binding
 spilloprotein A-IV
 RNA binding motif protein 10
 hypothetical protein DKFZp564K142 similar to implantation-associated protein
 lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase)
 Consensus includes gb:AF234255.1 /DEF=Homo sapiens clone KM36 immunoglobulin light chain variable region mRNA, partial cds. /FEA=mRNA /PROD=immunoglobulin lgi /AF234255
 gb:134698.1 /DEF=Homo sapiens T-cell receptor alpha (TCRA) mRNA (HLA-A3, 29, 87, 44, DR 2, 7), complete cds. /FEA=mRNA /GEN=TCRA /PROD=T-cell receptor alpha /134698
 hypothetical protein PP1057
 glucose-6-phosphatase, catalytic (glycogen storage disease type 1, von Gierke disease)
 protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 1
 transcriptional regulator interacting with the PHS-bromodomain 2
 peptidase (mitochondrial processing) beta
 enoyl Coenzyme A hydratase 1, peroxisomal
 promyelocytic leukemia
 KIAA1093 protein
 interferon gamma receptor 2 (interferon gamma transducer 1)
 solute carrier 18 (monocarboxylic acid transporters), member 8
 serine/threonine kinase 4
 IGF-II mRNA-binding protein 3
 aminopeptidase puromycin sensitive
 aldehyde dehydrogenase 2 family (mitochondrial)
 catenin (cadherin-associated protein), delta 1
 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 2 (8kD, B8)
 acrosomal vesicle protein 1
 Consensus includes gb:AA676803 /FEA=EST /DB_XREF=gi:2657325 /DB_XREF=est:j65504.s1 /CLONE=IMAGE:455119 /UG=Hs.13996 Homo sapiens cDNA, FLJ23260 fa, AA676803
 Consensus includes gb:AA676803 /FEA=EST /DB_XREF=gi:2657325 /DB_XREF=est:j65504.s1 /CLONE=IMAGE:455119 /UG=Hs.13996 Homo sapiens cDNA, FLJ23260 fa, AA676803
 phosphatidylinositol glycan, class H
 KIAA0630 protein
 GTPase regulator associated with the focal adhesion kinase pp125
 Consensus includes gb:AA927664 /FEA=EST /DB_XREF=gi:3076484 /DB_XREF=est:om71h10.s1 /CLONE=IMAGE:1552675 /UG=Hs.171626 transcription elongation factor B /AA927664
 glutaryl-Coenzyme A dehydrogenase
 hypothetical protein FLJ22578
 Consensus includes gb:AW517686 /FEA=EST /DB_XREF=gi:7155768 /DB_XREF=est:xs86405.x1 /CLONE=IMAGE:2776521 /UG=Hs.305923 ATPase, Ca++ transporting, plai /AW517686
 hypothetical protein MGC5306
 KIAA0904 protein
 acyl-Coenzyme A dehydrogenase, very long chain
 spleen tyrosine kinase
 Consensus includes gb:AU147402 /FEA=EST /DB_XREF=gi:11008923 /DB_XREF=est:AU147402 /CLONE=MAMMA1000567 /UG=Hs.287495 Homo sapiens cDNA FLJ12180 AU147402
 KIAA0515 protein
 phenylethanolamine N-methyltransferase
 GT198, complete ORF
 Consensus includes gb:BE328312 /FEA=EST /DB_XREF=gi:9202173 /DB_XREF=est:hs95412.x1 /CLONE=IMAGE:3144983 /UG=Hs.6614 ESTs, Weakly similar to A43932 m /BE328312
 artemin
 peroxisomal biogenesis factor 14
 hypothetical protein FLJ13491
 cytokine inducible SH2-containing protein
 Consensus includes gb:AF070580.1 /DEF=Homo sapiens clone 24527 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3387952 /UG=Hs.25422 Homo sapiens clone 24527 mRI /AF070580
 Cluster Ind. AL049370-Homo sapiens mRNA; cDNA DKFZp566D0918 (from clone DKFZp566D0918) /cds=UNKNOWN /gb=AL049370 /ig=4500162 /ug=Hs.13350 /len=2705
 zinc finger protein, X-linked
 putative capacitative calcium channel
 hypothetical protein FLJ13390 similar to PAF53
 zinc/fon regulated transporter-like
 tropomodulin
 mammin-binding lectin serine protease 1 (C4/C2 activating component of Ra-reactive factor)
 sbi-interactor 12 (SH3-containing protein)

culin 4B	AV694732	1,22
paired box gene 8	X69699	1,22
microfibrillar-associated protein 1	NM_005926	1,22
peroxisome biogenesis factor 10	BF339133	1,22
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 16	NM_003587	1,22
mitogen-activated protein kinase 9	U35002	1,22
Consensus includes gb:AK024988.1 /DEF=Homo sapiens cDNA: FLJ21335 fls, clone COL02546, /FEA=mRNA /DB_XREF=gi:10437419 /UG=Hs.303001 Homo sapiens cDNA: AK024988		1,22
enhancer of filamentation 1 (cas-like docking; CK-associated substrate related)	U84317	1,22
hypothetical protein FLJ13044	NM_024698	1,22
lymphoblastic leukemia derived sequence 1	BC002796	1,22
WD-repeat protein	BC001284	1,22
hypothetical protein FLJ22824	NM_024608	1,22
nuclear protein (KKEID repeat)	NM_006392	1,22
high-mobility group (nonhistone chromosomal) protein isoforms I and Y	AF176039	1,22
unit-related transcription factor 1 (acute myeloid leukemia 1; amt1 oncogene)	D43968	1,22
Consensus includes gb:AL120173 /FEA=EST /DB_XREF=gi:5926072 /DB_XREF=est:DKFZp781C117_s1 /CLONE=DKFZp781C117 /UG=Hs.301663 ESTs	AL120173	1,22
hypothetical protein FLJ20727	BE966019	1,22
stratifin	BC000329	1,22
growth hormone inducible transmembrane protein	AF131820	1,22
ret proto-oncogene (multiple endocrine neoplasia and medullary thyroid carcinoma 1, Hirschsprung disease)	M31213	1,22
nuclear protein family A, member 2 (HACA small nuclear RNPs)	BC000009	1,22
mitogen-activated protein kinase kinase kinase 1	BE646618	1,22
lectin, galactoside-binding, soluble, 3 (galectin 3)	BC001120	1,22
Consensus includes gb:A1949687 /FEA=EST /DB_XREF=gi:5741997 /DB_XREF=est:wt13g04.x1 /CLONE=IMAGE:2471190 /UG=Hs.173638 transcription factor 7-like 2 (T-cell)	AW006750	1,22
hypothetical protein FLJ20059	NM_005652	1,22
telomeric repeat binding factor 2	NM_005873	1,22
regulator of G-protein signalling 19	AV748469	1,22
SEC14-like 1 (S. cerevisiae)	NM_005930	1,22
meningioma expressed antigen 6 (coiled-coil proline-rich)	NM_004716	1,22
proprotein convertase subtilisin/kexin type 7	NM_004716	1,22
aminopeptidase puromycin sensitive	AW055008	1,22
H44 antigen	X91103	1,22
FGF receptor activating protein 1	AL049281	1,22
lysozyme homolog	NM_020426	1,22
tol-like receptor 7	NM_016562	1,22
dynactin intermediate chain 2	NM_023036	1,22
viral simian leukemia viral oncogene homolog B (ras related; GTP binding protein)	NM_002881	1,22
myotubularin related protein 3	AF233438	1,22
Consensus includes gb:AF184174 /DEF=Homo sapiens somatostatin receptor 2 (SSTR2) gene, complete cds, alternatively spliced /FEA=CDS_2 /DB_XREF=gi:7229402 /UG=Hs.184174		1,22
Consensus includes gb:AL034396 /DEF=Human DNA sequence from clone 1158812 on chromosome Xp11.21-11.4 Contains the ZXDA gene for X-linked duplicated Zinc finger	AF078844	1,22
RNA helicase-related protein	NM_017818	1,21
WD repeat domain 8	NM_024874	1,21
fibronectin 1	BC005858	1,21
heparan sulfate 2-O-sulfotransferase	AK023260	1,21
hypothetical protein FLJ10468	BC001651	1,21
HERV-H LTR-associated 2	NM_007072	1,21
hypothetical protein FLJ11413	NM_024554	1,21
solute carrier family 25 (mitochondrial carrier; oxoglutarate carrier), member 11	NM_003562	1,21
KIAA0210 gene product	A1300084	1,21
JMS protein	BC000484	1,21
hypothetical protein DKFZp434E2216	BC001043	1,21
GTT1 protein	NM_020151	1,21
RNA guanylyltransferase and 5'-phosphatase	AB012142	1,21
thymidine synthetase	NM_001071	1,21
Consensus includes gb:AL133099.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434B1521 (from clone DKFZp434B1521), /FEA=mRNA /DB_XREF=gi:6453555 /UG=Hs.27231	AL133099	1,21
Consensus includes gb:BE686922 /FEA=EST /DB_XREF=gi:10318698 /DB_XREF=est:601446588F1 /CLONE=IMAGE:3650432 /UG=Hs.161307 H3 histone, family 3A	BE686922	1,21
nuclear receptor subfamily 0, group B, member 1	NM_000475	1,21
stanniocalcin 2	BC000658	1,21
heat shock transcription factor 2	BC005329	1,21
topoisomerase (DNA) III alpha	A1878092	1,21
Consensus includes gb:AK024175.1 /DEF=Homo sapiens cDNA FLJ14113 fls, clone MAMMA1001715, /FEA=mRNA /DB_XREF=gi:10436491 /UG=Hs.323409 Homo sapiens	AK024175	1,21
heme-regulated initiation factor 2-alpha kinase	NM_014413	1,21

breast cancer metastasis-suppressor 1	AL050008	1,21
hypothetical protein RG083M05.2	AK026142	1,21
mitogen-activated protein kinase 7	U29725	1,21
G protein-coupled receptor 38	AL137000	1,21
81B and CNC homology 1, basic leucine zipper transcription factor 2	NM_021813	1,21
CD27-binding (Siva) protein	AW024335	1,21
uncharacterized hematopoietic stem/progenitor cells protein MDS028	NM_018463	1,21
NAD kinase	BE674658	1,21
PFTAIR protein kinase 1	AF119833	1,21
melanoma-associated antigen recognised by cytotoxic T lymphocytes	NM_006428	1,21
1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta)	U56418	1,21
KIAA1035 protein	NM_015239	1,21
F-box only protein 7	NM_012179	1,21
hypothetical protein DKFZp782A227	BC003163	1,21
Sle20-related serine/threonine kinase	NM_014720	1,21
KIAA0794 protein	BG339560	1,21
mitochondrial ribosomal protein L16	NM_017840	1,21
death effector domain-containing	AK022531	1,21
KIAA1449 protein	AB040882	1,21
Consensus includes gb:AI291720 /FEA=EST /DB_XREF=gi:3934494 /DB_XREF=est:cm80c05.x1 /CLONE=IMAGE:1855050 /UG=Hs.262642 ESTs	AI291720	1,21
hypothetical protein FLJ22551	NM_024708	1,21
heterogeneous nuclear ribonucleoprotein A1	AI144007	1,21
solute carrier family 25 (mitochondrial carrier, oxoglutarate carrier), member 11	AF070548	1,21
hypothetical protein FLJ20258	AF282167	1,21
neurocalcin delta	AF251081	1,21
GATA binding protein 3	AF796169	1,21
SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome)	AF100540	1,21
steroid regulatory element binding transcription factor 2	BE513151	1,21
oxidoreductase UCPA	NM_020139	1,21
hypothetical protein FLJ12171	NM_024619	1,21
a disintegrin and metalloproteinase domain 19 (meltrin beta)	Y13786	1,21
ATP-binding cassette, sub-family G (WHITE), member 5 (sterolin 1)	NM_022436	1,21
zinc finger protein 211	NM_066385	1,21
calcium channel, voltage-dependent, P/Q type, alpha 1A subunit	NM_023035	1,21
hypothetical protein	AC002550	1,21
KIAA1042 protein	A1633774	1,21
brain and reproductive organ-expressed (TNFRSF1A modulator)	U19178	1,21
cyclin G1	BC000196	1,21
arylalkylamine N-acetyltransferase	NM_001088	1,21
KIAA0508 protein	H92070	1,21
cyclin-dependent kinase 2	AB012305	1,21
ribosomal protein L5	BF214492	1,21
hypothetical protein 628	BC000974	1,21
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6 (17kD, B17)	NM_002493	1,21
dipeptidylpeptidase 8	NM_017743	1,21
hypothetical protein	NM_013313	1,21
cytochrome c oxidase subunit VIc	NM_001867	1,21
hypothetical protein RG083M05.2	AI337901	1,21
a disintegrin and metalloproteinase domain 22	NM_021721	1,21
guanine nucleotide binding protein (G protein), gamma 7	NM_005145	1,21
diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)	NM_020548	1,21
tropomyosin 4	AI214061	1,21
Consensus includes gb:AK025152.1 /DEF=Homo sapiens cDNA: FLJ21499 fls, clone COL05634 /FEA=mRNA /DB_XREF=gi:10437612 /UG=Hs.306773 Homo sapiens cDNA: AK025152	NM_014753	1,21
KIAA0187 gene product	AW340333	1,21
IKK-related kinase epsilon; inducible IkappaB kinase	NM_002557	1,21
oviductal glycoprotein 1, 120kD (mucin 9, oviductin)	W60802	1,21
hypothetical protein FLJ12438	AU149503	1,21
Ras-GTPase activating protein SH3 domain-binding protein 2	BC004170	1,21
polymerase (DNA directed), epsilon 3 (p17 subunit)	AF275803	1,21
Slah-interacting protein	AA565715	1,21
neuretic peptide receptor B/guanlylate cyclase B (atriatriuretic peptide receptor B)	NM_014786	1,21
KIAA0337 gene product	NM_017657	1,21
hypothetical protein FLJ20080		

Consensus includes gb:N36842 /FEA=EST /DB_XREF=gi:1157984 /DB_XREF=gi:1157984 /JG=Hs.274412 similar to yeast Upf3, variant A	1,21
KIAA00431 protein	1,21
TAR (HIV) RNA binding protein 2	NM_004178
hypothetical protein FLJ20169	NM_017704
polynucleotide kinase 3'-phosphatase	NM_007254
chromosome 8 open reading frame 9	NM_022107
hypothetical protein FLJ20123	NM_017674
origin recognition complex, subunit 4-like (yeast)	AF047598
hypothetical protein MGC4172	NM_024308
hypothetical protein FLJ23018	NM_024810
KIAA1139 protein	AF070096
peroxisomal biogenesis factor 11B	NM_003846
choline kinase	NM_001277
Consensus includes gb:U02632.1 /DEF=Human calcium-activated potassium channel mRNA, partial cds. /FEA=mRNA /PROD=calcium-activated potassium channel /DB_XREF=U02632	BE969622
synaptin 3A	BF508616
KIAA0211 gene product	AA845258
biglycan	D50740
thyroid transcription factor 1	NM_024030
SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome)	AF100539
neurogranin (protein kinase C substrate, RC3)	NM_003178
Consensus includes gb:AL042729 /FEA=EST /DB_XREF=gi:5935545 /DB_XREF=est:DKFZp434B0222_s1 /CLONE=DKFZp434B0222 JG=Hs.112023 ESTs	AL042729
Consensus includes gb:RZ7378 /FEA=EST /DB_XREF=gi:7835113 /DB_XREF=est:V53611.1 /CLONE=IMAGE:133508 JG=Hs.322426 ESTs, Moderately similar to TIAR_HUI	RZ7378
Consensus includes gb:U38276 /FEA=EST /DB_XREF=gi:5935545 /DB_XREF=est:V53611.1 /CLONE=IMAGE:133508 JG=Hs.322426 ESTs, Moderately similar to TIAR_HUI	U38276
semA domain, immunoglobulin domain (lig), short basic domain, secreted, (semaphorin) 3F	BE259395
small nuclear RNA activating complex, polypeptide 5, 19KD	NM_003529
H3 histone family, member A	NM_003645
faty-acid-Coenzyme A ligase, very long-chain 1	AF080800
hypothetical protein DKFZ0564D0478	AF339834
Consensus includes gb:A554879 /FEA=EST /DB_XREF=gi:4487242 /DB_XREF=est:ie53603.x1 /CLONE=IMAGE:2090428 /JG=Hs.81360 Homo sapiens cDNA: FLJ21927 fls	AF339834
karyopherin (importin) beta 3	AF080129
DKFZP434D193 protein	NM_003433
zinc finger protein 132 (clone pT2-12)	NM_015165
Cdc42 guanine exchange factor (GEF) 9	AF123652
leucine zipper, putative tumor suppressor 1	AF199061
fem-1 homolog b (C. elegans)	NM_004852
one cut domain, family member 2	NM_001827
CDC28 protein kinase 2	BC491844
v-jun sarcoma virus 17 oncogene homolog (avian)	AF197627
solute carrier family 19 (folate transporter), member 1	NM_003730
conserved gene amplified in osteosarcoma	NM_007429
Cluster int. AN014299:U1-H-B109-aax-a-07-Q-U1.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:2710573 /clone_end=3' /gb=AW014299 /igf=5863056 /JG=Hs.238968 /len=391	AF080579
Consensus includes gb:AL134724 /FEA=EST /DB_XREF=gi:6602911 /DB_XREF=est:DKFZp547P246_s1 /CLONE=DKFZp547P246 JG=Hs.4988 Homo sapiens clone 24711	AF080579
Consensus includes gb:AL134724 /FEA=EST /DB_XREF=gi:6602911 /DB_XREF=est:DKFZp547P246_s1 /CLONE=DKFZp547P246 JG=Hs.4988 Homo sapiens clone 24711	M85836
succinate dehydrogenase complex, subunit C, integral membrane protein, 15KD	AF080579
v-akt murine thymoma viral oncogene homolog 2	AF080579
brromodomain-containing 4	AF080579
myotubularin related protein 4	AF080579
calcium/calmodulin-dependent protein kinase 1	AF080579
ChpP caseinolytic protease, ATP-dependent, proteolytic subunit homolog (E. coli)	AF080579
chronic somatomammotropin hormone 1 (placental lactogen)	AF080579
nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon	AF080579
KIAA0445 gene product	AF080579
steroid O-acetyltransferase (acyl-Coenzyme A, cholesterol acyltransferase) 1	AF080579
formin-like	AF080579
protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcinurin A gamma)	AF080579
tumor necrosis factor receptor superfamily, member 5	AF080579
CG-112 protein	AF080579
ubiquitin specific protease 7, paraplegin (pure and complicated autosomal recessive)	AF080579
ubiquitin specific protease 25	AF080579
Rho-associated, coiled-coil containing protein kinase 2	AF080579
hypothetical protein FLJ11011	AF080579
hypothetical protein FLJ10737	AF080579
KIAA0592 protein	AF080579
hypothetical protein FLJ10486	AF080579

cold inducible RNA binding protein
 super conserved receptor expressed in brain 3
 fibronectin 1
 LM protein (similar to rat protein kinase C-binding enzyme)
 glycoprotein V (platelet)
 Consensus includes gb:AU134977 /FEA=EST /DB_XREF=gi:10995516 /DB_XREF=est:AU134977 /CLONE=PLACE100926 /UG=Hs.322149 Human clone 137308 mRNA, par
 prefoldin 1
 rab6 GTPase activating protein (GAP and centrosome-associated)
 olfactory receptor, family 2, subfamily F, member 1
 phospholipid acid phosphatase type 2B
 KIAA0438 gene product
 KIAA0792 gene product
 putative L-type neutral amino acid transporter
 extra spindle poles, S. cerevisiae, homolog of
 Xq28. 2000bp sequence contig. ORF
 hypothetical protein FLJ10597
 CDC28 protein kinase 1
 hypothetical protein FLJ20399
 COP9 (constitutive photomorphogenic, Arabidopsis, homolog) subunit 7B
 endothelin receptor type A
 hypothetical protein FLJ11289
 solute carrier family 29 (nucleoside transporters), member 2
 zinc finger protein-like 1
 DEAD/HD (Asp-Glu-Ala-Asp/His) box polypeptide 34
 phospholipase C, epsilon 2
 nitrogen fixation cluster-like
 solute carrier family 1 (glutamate/neutral amino acid transporter), member 4
 FKBP58 binding protein 8 (38kD)
 DKE2P434F195 protein
 sulfotransferase family, cytosolic, 1A, phenol-preferring, member 2
 enhancer of zeste homolog 1 (Drosophila)
 M-phase phosphoprotein 9
 protein tyrosine phosphatase, receptor type, F
 oxidative-stress responsive 1
 sudD (suppressor of bimD6, Aspergillus nidulans) homolog
 6.2 kd protein
 neuron-specific protein
 hypothetical protein FLJ20125
 Consensus includes gb:AW293356 /FEA=EST /DB_XREF=gi:6699992 /DB_XREF=est:LIU-H-B12-ahl-c-11-Q-UI.s1 /CLONE=IMAGE:2727165 /UG=Hs.58220 Homo sapiens cDN
 selenophosphate synthetase 2
 inositol(myo)-1(or 4)-monophosphatase 1
 seven transmembrane domain orphan receptor
 gb:L14454.1 /DEF=Human Ig rearranged mu-chain gene V-N-D-N-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398495 /FL=gb:L14454.1
 KIAA0467 protein
 tumor protein D52-like 2
 uncharacterized bone marrow protein BM039
 zinc finger protein 282
 small inducible cytokine subfamily B (Cys-X-Cys), member 6 (granulocyte chemolactic protein 2)
 Consensus includes gb:AL109705.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 73337. /FEA=mRNA /DB_XREF=gi:5689834 /UG=Hs.9997 Homo
 crooked neck protein (cm)
 Consensus includes gb:AA001552 /FEA=EST /DB_XREF=gi:1437017 /DB_XREF=est:ze46c08.s1 /CLONE=IMAGE:362030 /UG=Hs.59839 ESTs, Weakly similar to CGH7L c
 insulin induced protein 2
 hypothetical protein FLJ20259
 hypothetical protein FLJ10656
 huntingtin (Huntington disease)
 hypothetical protein FLJ12565
 granuln
 golgi SNAP receptor complex member 2
 hypothetical protein FLJ21865
 I-complex 10 (a murine tap homolog)
 mitogen-activated protein kinase 2
 KIAA0650 protein
 male-specific lethal 3-like 1 (Drosophila)

protein phosphatase 2, regulatory subunit B (B56), gamma isoform	L42375	1,20
CGI-92 protein	NM_016035	1,20
polymerase (RNA) II (DNA directed) polypeptide A (220kD)	NM_000937	1,20
H2A histone family, member G	NM_021065	1,20
retinoblastoma-like 2 (p130)	BF110947	1,20
calcineurin binding protein 1	AB002328	1,20
HIV-1 Tat interactive protein 2, 30 kDa	BC002439	1,20
phosphoribosylformylglycinamide synthase (FGAR amidotransferase)	AL044328	1,20
UDP glycosyltransferase 1 family, polypeptide A6	NM_001072	1,20
persephin	NM_004158	1,20
hypothetical protein FLJ20298	NM_017750	1,20
linked to Surfeit genes in Fugu rubripes 2; LSFR2 gene 2	AI684664	1,20
sodium carrier family 2 (facilitated glucose transporter), member 2	NM_000340	1,20
A kinase (PRKA) anchor protein 1	U34074	1,20
hypothetical protein FLJ12553	NM_024700	1,20
KIAA0842 protein	AB020849	1,20
zinc finger protein, Y-linked	NM_003411	1,20
KIAA1254 protein	AB033080	1,20
HpaII lny fragments locus 9C	NM_022727	1,20
hypothetical protein FLJ20344	NM_017776	1,20
synaptosomal-associated protein, 23kD	U55936	1,20
inositol polyphosphate-5-phosphatase, 75kD	AK022846	1,20
aconitase 1, soluble	NM_002197	1,20
peroxisomal D3,D2-enoyl-CoA isomerase	NM_006117	1,20
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 5 (13kD, B13)	AK022209	1,20
phospholipid scramblase 3	AI783924	1,20
nigamap homolog 1 (C. elegans)	AW063371	1,20
nucleic (nucleoside diphosphate linked moiety X)-type motif 9	NM_024047	1,20
KIAA0047 gene product	BE865244	1,20
hypothetical protein FLJ20113	AL098714	1,20
E74-like factor 2 (ets domain transcription factor)	AF286223	1,20
suppressor of Ty 6 homolog (S. cerevisiae)	AI494567	1,20
mitogen-activated protein kinase 14	AF100544	1,20
KIAA0244 protein	BF430956	1,20
cathepsin D (lysosomal aspartyl protease)	NM_001909	1,20
chromosome 18 open reading frame 1	NM_004338	1,20
synovial sarcoma, X breakpoint 1	BC001003	1,20
hypothetical protein FLJ20477	NM_017837	1,20
WD repeat domain 13	AF158978	1,19
hypothetical protein DKFZp762P2111	AU151157	1,19
amiloride-sensitive cation channel 3, testis	NM_020322	1,19
DKFZP434D1335 protein	BC054966	1,19
entyrin repeat domain 5	NM_022096	1,19
HSPC052 protein	NM_014150	1,19
mucosa associated lymphoid tissue lymphoma translocation gene 1	AB026118	1,19
glutathione reductase	NM_000637	1,19
GMY2 ganglioside activator protein	X61094	1,19
KIAA0191 protein	AI049862	1,19
ubimucdin 1	NM_016938	1,19
Consensus includes gb:AW517716 /FEA=EST /DB_XREF=gi:7155798 /DB_XREF=est:xa86h03.x1 /CLONE=IMAGE:2776565 /UG=Hs.293472 ESTs	AW517716	1,19
splicing factor (CC1.3)	AI690947	1,19
zinc finger protein 36, C3H type, homolog (mouse)	NM_003407	1,19
KIAA0440 protein	NM_015556	1,19
glycoprotein A repetitions predominant	NM_005512	1,19
crystallin, alpha A	AP001748	1,19
protocadherin LKC	NM_017675	1,19
adenosine deaminase, RNA-specific	NM_001111	1,19
regulator of nonsense transcripts 1	D86988	1,19
sacrospinal vesicle protein 1	NM_001612	1,19
CGI-96 protein	Z93241	1,19
pleomorphic adenoma gene-like 2	NM_002857	1,19
hypothetical protein FLJ22418	NM_024626	1,19
v-yes-1 Yamaguchi sarcoma viral related oncogene homolog	NM_002350	1,19

8100 calcium binding protein A13	NM_005878	1.19
peroxisomal fatty-acylated protein	NM_002857	1.19
hypothetical protein FLJ10120	NM_018001	1.19
pasillin	NM_019049	1.19
nuclear receptor coactivator 2	NM_002859	1.19
guanine nucleotide binding protein (G protein), alpha z polypeptide	NM_006540	1.19
ribosomal protein L26	NM_002073	1.19
BTAF1 RNA polymerase II, B-TFIIID transcription factor-associated, 170 kD (Maf1 homolog, S. cerevisiae)	AL121871	1.19
guanine nucleotide binding protein 10	A001017	1.19
Consensus includes gb-AA156865 /FEA=EST /DB_XREF=gi:1728480 /DB_XREF=est:20108.s1 /CLONE=IMAGE:502503 /UG=Hs.22370 Homo sapiens mRNA: cDNA DKFZp	NM_004125	1.19
protein tyrosine phosphatase, receptor type, C-associated protein	AA156865	1.19
FGF receptor activating protein 1	NM_005608	1.19
Wiskott-Aldrich syndrome (eczema-thrombocytopenia)	AL049261	1.19
lysine 3-monooxygenase/hypophan 5-monooxygenase activation protein, theta polypeptide	U12707	1.19
protein kinase C, zeta	AA854017	1.19
elastase 3B, pancreatic	NM_002744	1.19
urothrin (homologous to dystrophin)	NM_007352	1.19
serine/threonine kinase 25 (STE20 homolog, yeast)	NM_007124	1.19
synthrophin, beta 2 (dystrophin-associated protein A1, 59kD, basic component 2)	NM_006374	1.19
chromobox homolog 3 (HP1 gamma homolog, Drosophila)	AA741303	1.19
thyroid hormone receptor interactor 7	NM_016587	1.19
chromobox homolog 6	AF274949	1.19
I-kappa-B-interacting Ras-like protein 2	NM_014292	1.19
KIA0687 protein	NM_017595	1.19
bridging integrator 3	AB66295	1.19
hypothetical protein MGC3162	AK001289	1.19
calcium channel, voltage-dependent, gamma subunit 2	NM_024078	1.19
signal sequence receptor, alpha (translocin-associated protein alpha)	AL022313	1.19
CD209 antigen	AW006345	1.19
hypothetical protein PRO0943	NM_021155	1.19
BCL2/adenovirus E1B 19kD interacting protein 3	NM_018568	1.19
muskelin 1, intracellular mediator containing kelch motifs	NM_004052	1.19
G protein-binding protein CRFG	NM_013255	1.19
hypothetical protein FLJ13117	A081107	1.19
DKFZP586J1624 protein	NM_023071	1.19
hypothetical protein FLJ21212	NM_015537	1.19
cyclin M4	NM_024642	1.19
Consensus includes gb-AA885297 /FEA=EST /DB_XREF=gi:2994374 /DB_XREF=est:al58n03.s1 /CLONE=IMAGE:1461557 /UG=Hs.323567 CD36 antigen (collagen type I rec	NM_020184	1.19
protein homologous to salivary proline-rich protein P-8	AA885297	1.19
GDNF family receptor alpha 2	NM_012390	1.19
solute carrier family 24 (sodium/potassium/calcium exchanger), member 3	U97145	1.19
protein tyrosine phosphatase type IVA, member 1	NM_020689	1.19
zinc finger protein 174	BF576710	1.19
hypothetical protein FLJ11128	BC001161	1.19
heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kD)	NM_018332	1.19
KIA09523 protein	M84630	1.19
cell matrix adhesion regulator	AU152563	1.19
chromosome 8 open reading frame 2	NM_005200	1.19
Consensus includes gb-AK024450.1 /DEF=Homo sapiens mRNA for FLJ00042 protein, partial cds. /FEA=mRNA /GEN=FLJ00042 protein /DB_XREF=gi:104 AK024450	T90773	1.19
copine III	AA541758	1.19
Consensus includes gb-AF228166.1 /DEF=Homo sapiens docking protein 1-like protein mRNA, partial cds. /FEA=mRNA /PROD=docking protein 1-like protein /DB_XREF=gi:71 AF229166	AF229166	1.19
small nuclear ribonucleoprotein polypeptide C	NM_003093	1.19
PR domain containing 5	NM_018699	1.19
glycine C-acetyltransferase (2-amino-3-ketobutyrate coenzyme A ligase)	NM_014281	1.19
nuclear autoantigenic sperm protein (histone-binding)	AY003362	1.19
Arg/Arg-interacting protein ArgBP2	NM_021069	1.19
muscleblind-like (Drosophila)	NM_021038	1.19
hypothetical protein FLJ13433	NM_022486	1.19
hypothetical protein FLJ10900	NM_018284	1.19
methyl-CpG binding domain protein 1	NM_015946	1.19
leukotriene A4 hydrolase	J02959	1.19
hypothetical protein FLJ20241	NM_017721	1.19

N-acetylserine amidohydrolase (acid ceramidase)-like
dendritic cell protein
RAD1 homolog (S. pombe)
serine/threonine kinase 18
tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory)
hypothetical protein FLJ20160
HIV-1 Tat interactive protein, 60 kDa
hypothetical protein FLJ10848
fibronectin type 3 and SPRY domain-containing protein
Fas (TNFRSF6) associated factor 1
hypothetical protein dJ472B4
hypothetical protein FLJ10706
RacO protein-like (DNA helicase O1-like)
hypothetical protein MGC10772
KIAA0874 protein
cytochrome c oxidase subunit VIc pseudogene 1
kraken-like
Consensus includes gb:AL050035.1 /DEF=Homo sapiens mRNA; cDNA DKFp566H0124 (from clone DKFp566H0124) /FEA=mRNA /DB_XREF=gi:4884276 /UG=Hs.13313
core promoter element binding protein
cadherin 6, type 2, K-cadherin (fetal kidney)
hypothetical protein FLJ10081
TPA regulated locus
somatostatin receptor 1
kallinin p80 (WD40-containing) subunit B 1
follicle stimulating hormone receptor
high-mobility group 20A
I-box and leucine-rich repeat protein 4
SRY (sex determining region Y)-box 1
chemokine-like receptor 1
Consensus includes gb:AL117381 /DEF=Human DNA sequence from clone RPS-457M17 on chromosome 20 Contains ESTs, STSs, GSSs and CpG islands, Contains the 3' pA
RNA cyclase homolog
transient receptor potential channel 1
KIAA0056 protein
chromosome 21 open reading frame 45
replication protein A3 (14kD)
ATPase, H⁺ transporting, lysosomal (vacuolar proton pump), alpha polypeptide, 70kD, isoform 1
KIAA0978 protein
similar to hypothetical protein MNCb-2366
tripartite motif-containing 31
cotomer protein complex, subunit beta
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 (GalNAc-T2)
cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)
Notch homolog 3 (Drosophila)
caspase 9, apoptosis-related cysteine protease
guanine nucleotide binding protein (G protein), beta polypeptide 1
Consensus includes gb:AU148611 /FEA=EST /DB_XREF=gi:11010132 /DB_XREF=est:AU148611 /CLONE=NT2RM400607 /UG=Hs.296832 Human pTR7 mRNA for repeliv
PTD002 protein
gb:BC001788.1 /DEF=Homo sapiens, Similar to hypothetical protein from EUROMAGE 42353, clone MGC:2806, mRNA, complete cds. /FEA=mRNA /PROD=Similar to hypoth
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 1 (GalNAc-T1)
Deleted in split-hand/split-foot 1 region
acetylserotonin O-methyltransferase-like
KIAA0100 gene product
Consensus includes gb:AV701318 /FEA=EST /DB_XREF=gi:10717648 /DB_XREF=est:AV701318 /CLONE=ADAAGD10 /UG=Hs.276626 ArgAbi-interacting protein ArgBP2
CGL-87 protein
Nileperch breakage syndrome 1 (nibrin)
JMI1 protein
hypothetical protein FLJ12785
maternally expressed 3
zinc finger protein 208
gb:NM_000256.2 /DEF=Homo sapiens myosin-binding protein C, cardiac /DB_XREF=gi:13325073 /PROD=protein C, cardiac /DB_XREF=gi:13325073
menage a trois 1 (CAK assembly factor)
aldo-keto reductase family 1, member C2 (dihydrodiol dehydrogenase 2; bile acid binding protein; 3-alpha hydroxysteroid dehydrogenase, type III)

AV724215 1.19
NM_006360 1.19
AF084513 1.19
AB020739 1.19
BF347089 1.19
NM_017694 1.19
AA886971 1.19
NM_018243 1.19
NM_030800 1.19
NM_024333 1.19
NM_007051 1.19
NM_019556 1.19
NM_018186 1.19
AI655944 1.19
NM_030567 1.19
AB020681 1.19
AF042163 1.19
AL157851 1.19
AL050035 1.19
AB017493 1.19
BC000019 1.19
NM_017981 1.19
AF764378 1.19
NM_001049 1.19
NM_005886 1.19
M95489 1.19
NM_018200 1.19
AF176699 1.19
NM_005986 1.18
U75526 1.18
NM_005772 1.18
NM_003304 1.18
AI796581 1.18
NM_018944 1.18
BC005264 1.18
NM_001690 1.18
M64760 1.18
BG230758 1.18
AF230386 1.18
NM_016451 1.18
NM_004481 1.18
AI871089 1.18
AW139152 1.18
AB015653 1.18
NM_002074 1.18
NM_014861 1.18
NM_016144 1.18
AV692127 1.18
NM_006304 1.18
AA689799 1.18
AA904674 1.18
AV701318 1.18
NM_016030 1.18
AK001017 1.18
NM_014008 1.18
NM_024655 1.18
AF118863 1.18
NM_007153 1.18
NM_000256 1.18
NM_002431 1.18
U05598 1.18

mevalonate kinase (mevalonic aciduria) 1.18 M85488
 hypothetical protein FLJ10604 1.18 NM_018154
 TNF receptor-associated factor 4 1.18 AF082185
 vesicle-associated membrane protein 3 (collubrevin) 1.18 BC005941
 protein tyrosine phosphatase, receptor type 1 (polypeptide) (PTPRF), interacting protein (liprin), alpha 1 1.18 A4195259
 synaptobin, beta 1 (synaptobin-associated protein A1, 59kD, basic component 1) 1.18 BG464314
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1 1.18 A1554300
 HCDI protein 1.18 A1347090
 trophobin 1.18 UD4811
 chloride channel 3 1.18 NM_001829
 ubiquitin specific protease 11 1.18 BC000350
 Consensus includes gb:A0971258 /FEA=EST /DB_XREF=gi:5768084 /DB_XREF=est:wr27b08.x1 /CLONE=IMAGE:2488887 /UG=Hs.295923 seven in absentia (Drosophila) ho A1971258 1.18
 solute carrier family 20 (phosphate transporter), member 2 1.18 NM_006749
 F-box only protein 4 1.18 NM_018007
 Consensus includes gb:AL359931.1 /DEF=Homo sapiens mRNA full length insert cDNA done EUROMIMAGE 288936 /FEA=mRNA /DB_XREF=gi:8977886 /UG=Hs.48607 Hrr AL359931 1.18
 high-mobility group (nonhistone chromosomal) protein 2 1.18 BC006903
 KIAA1240 protein 1.18 AB033066
 ribosomal protein L23a 1.18 NM_000984
 3-hydroxyisobutyryl-Coenzyme A hydrolase 1.18 AWO00964
 LIM domain kinase 1 1.18 AF134379
 RNA binding protein S1, serine-rich domain 1.18 NM_006711
 hypothetical protein MGC3035 1.18 AL040896
 nuclear transcription factor Y, gamma 1.18 U78774
 junctional adhesion molecule 1 1.18 AF154005
 myosin, light polypeptide 4, alkali: atrial, embryonic 1.18 X52005
 cholinergic receptor, nicotinic, epsilon polypeptide 1.18 A1850860
 hypothetical protein FLJ21047 1.18 NM_024569
 ADP-ribosylation factor GTPase activating protein 1 1.18 BC005122
 Consensus includes gb:AW094068 /FEA=EST /DB_XREF=gi:6039220 /DB_XREF=est:xc26c06.x1 /CLONE=IMAGE:2585386 /UG=Hs.110630 Human BRCA2 region. mRNA s-AW094068 1.18
 hypothetical protein FLJ11028 1.18 NM_018303
 flightless 1 homolog (Drosophila) 1.18 BG421186
 gene predicted from cDNA with a complete coding sequence 1.18 NM_014628
 mitochondrial translational initiation factor 2 1.18 NM_002453
 G protein-coupled receptor kinase-interactor 1 1.18 NM_014030
 melanin 1 1.18 BC001906
 N-ethylmaleimide-sensitive factor 1.18 NM_008178
 hepatitis B virus x-interacting protein (8.6kD) 1.18 NM_008402
 cell division cycle 2-like 5 (cholinesterase-related cell division controller) 1.18 NM_003718
 cardiac-specific homeo box 1.18 NM_004387
 retinol binding protein 4, plasma 1.18 NM_008744
 gastric intrinsic factor (vitamin B synthesis) 1.18 NM_005142
 DiGeorge syndrome critical region gene DGS1; likely ortholog of mouse expressed sequence 2 embryonic lethal 1.18 NM_022719
 hypothetical protein MGC10682 1.18 BC004952
 hypothetical protein FLJ21413 1.18 NM_024738
 TATA-binding protein-binding protein 1.18 NM_013375
 Consensus includes gb:A052538 /FEA=EST /DB_XREF=gi:3308527 /DB_XREF=est:oz27g09.x1 /CLONE=IMAGE:1676608 /UG=Hs.31834 Homo sapiens clone 25129 mRNA A052538 1.18
 hypothetical protein FLJ13046 similar to exorphin 4 1.18 NM_022459
 peroxisomal biogenesis factor 14 1.18 AB017546
 capping protein (actin filament), gelsolin-like 1.18 NM_001747
 cell division cycle 2-like 5 (cholinesterase-related cell division controller) 1.18 BE875449
 KIAA0173 gene product 1.18 AL043927
 Consensus includes gb:AL008730 /DEF=Human DNA sequence from PAC 487J7 on chromosome 6q21.22.1. Contains an unknown gene coding for three alternative mRNAs. 'AL008730 1.18
 protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha) 1.18 AL353950
 oxidation resistance 1 1.18 NM_018002
 hypothetical protein FLJ20195 1.18 NM_017706
 KIAA0453 protein 1.18 BF111982
 hypothetical protein FLJ13153 1.18 AF267659
 Rho GTPase activating protein 8 1.18 NM_015366
 hypothetical protein FLJ13348 1.18 NM_024918
 gb:AF271775.1 /DEF=Homo sapiens DC48 mRNA, complete cds. /FEA=mRNA /PROD=DC48 mRNA, complete AF271775 1.18
 tachykinin 3 (neuromedin K, neurokinin beta) 1.18 NM_013251
 timeless homolog (Drosophila) 1.18 NM_003920
 KIAA0682 gene product 1.18 NM_016196

FIGURE 4B.

gene	NCBI Accession Number	FOLD DECREASE
Consensus includes gb:AA292281 /FEA=EST /DB_XREF=gi:1940261 /DB_XREF=est:251003.s1 /CLONE=IMAGE:725837 /UG=Hs.181307 H3 histone, family 3A	AA292281	15.15
carotidin	AA292281	13.38
ribosomal protein S11	AF187078	10.86
ribosomal protein S19	BF680255	9.95
ribosomal protein, large P2	BC000023	9.72
Consensus includes gb:AW302047 /FEA=EST /DB_XREF=gi:5711724 /DB_XREF=est:x52108.x1 /CLONE=IMAGE:2763783 /UG=Hs.76230 ribosomal protein S10	AW302047	8.60
Consensus includes gb:LA48784 /DEF=050 Homo sapiens cDNA /FEA=mRNA /DB_XREF=gi:1066715 /UG=Hs.182426 ribosomal protein S2	BC005354	7.66
Consensus includes gb:LA48784 /DEF=050 Homo sapiens cDNA /FEA=mRNA /DB_XREF=gi:1066715 /UG=Hs.182426 ribosomal protein S2	LA48784	7.66
Consensus includes gb:AJ249377.1 /DEF=Homo sapiens partial mRNA for human Ig lambda light chain variable region, clone MB91 (331 bp). /FEA=mRNA (GENE=IGLV /P /J249377	AW302047	7.66
Consensus includes gb:D84143.1 /DEF=Human immunoglobulin (mAb59) light chain V region mRNA, partial sequence. /FEA=mRNA /PROD=immunoglobulin light chain V D84143	BE737027	6.93
ribosomal protein L27a	BE737027	6.41
emopamil binding protein (steroid isomerase)	N58483	5.40
carotidin	AA910371	5.23
hypothetical protein PRO1843	NM_018507	4.91
Cluster Ind. A1201594:gc02h12.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:1708487 /clone_end=3 /gb=A1201594 /gi=3754200 /ug=Hs.239333 /len=591	A1201594	4.69
ribosomal protein S20	AF113008	4.57
C-terminal binding protein 1	BF984434	4.45
KIA00908 protein	AA908765	4.42
ribosomal protein L27	BE312027	4.24
Consensus includes gb:AF044592 /DEF=Homo sapiens lymphocyte-predominant Hodgkins disease case 4 immunoglobulin heavy chain gene, variable region, partial cds // AF044592	NM_024940	4.13
hypothetical protein FLJ21034	BM153399	4.04
aminopeptidase puromycin sensitive	AF043586	4.00
immunoglobulin lambda locus	AJ239383	3.96
Consensus includes gb:AJ239383.1 /DEF=Homo sapiens mRNA for immunoglobulin heavy chain variable region, ID 31. /FEA=mRNA (GENE=IGHV /PROD=immunoglobulin	AJ239383	3.81
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2	X93598	3.78
hypothetical protein FLJ12619	BE465032	3.67
lymphoid blast crisis oncogene	AF127481	3.63
Consensus includes gb:AV719355 /FEA=EST /DB_XREF=gi:10816507 /DB_XREF=est:AV719355 /CLONE=GLCEMB08 /UG=Hs.97109 ESTs	AV719355	3.59
alanine-glyoxylate aminotransferase 2-like 1	NM_031279	3.56
Cluster Ind. A1949010:wg36807.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2473332 /clone_end=3 /gb=A1949010 /gi=5741320 /ug=Hs.104036 /len=457	A1949010	3.49
PCTAIRE protein kinase 1	NM_006201	3.42
ADP-ribosylation factor 6	AA502643	3.42
tyrosine 3-monooxygenase/tyriophan 5-monooxygenase activation protein, epsilon polypeptide	AA502643	3.37
Consensus includes gb:BF973387 /FEA=EST /DB_XREF=gi:12340602 /DB_XREF=est:60224233F1 /CLONE=IMAGE:4330861 /UG=Hs.305989 Human DNA sequence // BF973387	BF973387	3.33
Q protein-coupled receptor kinase 6	BC423052	3.24
M10098 Human 18S rRNA sequence, length 1968 bases, middle target bases 647-1292	M10098	3.24
GM2 ganglioside activator protein	X61094	3.24
dystrophin myotonic-containing WD repeat motif	L19267	3.24
collagen, type I, alpha 1	A1743521	3.22
cathepsin S	BC026842	3.19
transition initiation factor IF2	AB016284	3.16
hypothetical protein FLJ22965	NM_022101	3.14
coactivator-associated arginine methyltransferase-1	AL529398	3.13
JTV1 gene	AF116615	3.10
RA86 Interacting, kinesin-like (rakinesin 6)	NM_005733	3.06
hypothetical protein FLJ20666	NM_018333	3.04
isodihydrodehydrogenase 1 (NADP+), soluble	NM_005896	3.03
Consensus includes gb:AI524687 /FEA=EST /DB_XREF=gi:4438822 /DB_XREF=est:h12a07.x1 /CLONE=IMAGE:2118036 /UG=Hs.57969 phenylalanine-tRNA synthetase A1524687	AI524687	3.03
Consensus includes gb:AL138178 /DEF=Human DNA sequence from clone RP3-322L4 on chromosome 6. Contains the SOX4 gene for SRY (sex determining region Y)-box AL138179	NM_003787	3.02
nucleolar protein 4	AW303136	2.98
ribosomal protein L38	BC000603	2.91
ribosomal protein L38	BC000603	2.90
Consensus includes gb:AW090043 /FEA=EST /DB_XREF=gi:6047387 /DB_XREF=est:h01c05.x1 /CLONE=IMAGE:2592488 /UG=Hs.326464 Homo sapiens cDNA: FLJ21 AW090043	AW090043	2.88
Consensus includes gb:AW971415 /FEA=EST /DB_XREF=gi:8161260 /DB_XREF=est:EST338304 /UG=Hs.165337 ESTs	AW971415	2.87
J04423 E coli bioB gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	J04423	2.85
Consensus includes gb:AA890010 /FEA=EST /DB_XREF=gi:3016889 /DB_XREF=est:ai8908.s1 /CLONE=IMAGE:1403679 /UG=Hs.50785 SEC22, vesicle trafficking protol AA890010	AA890010	2.83
gb:Z25437.1 /DEF=H sapiens protein-tyrosine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-tyrosine kinase /DB_XREF=gi:405752 /FL=gb:Z25437.1	Z25437	2.81
preludin 4	NM_002823	2.85
KIA11046 protein	NM_014928	2.83
ATPase, H+ transporting, lysosomal (vacuolar proton pump) non-catalytic accessory protein 1A (110/116KD)	NM_005177	2.81

hypothetical protein FLJ10159 NM_018013 2.80
Consensus includes gb:BG109746 /FEA=EST /DB_XREF=gi:12603252 /DB_XREF=est:602260883F1 /CLONE=IMAGE:4366381 /UG=Hs.325625 Homo sapiens clone 239 BG109746 2.79
HIV-1 rev binding protein 2 A912583 2.76
acidic epidermal glycoprotein-like 1 X95238 2.75
hypothetical protein FLJ12619 AL136632 2.74
hypothetical protein FLJ14107 NM_025026 2.73
Escherichia coli /REF=J04423 /DEF=E coli bioB gene biotin synthetase corresponding to nucleotides 2393-2682 of J04423 /LEN=1114 (-5'..-M..-3' represent transcript region) 2.73
E74-like factor 4 (ets domain transcription factor) J04423 2.71
ribonuclease P, 40KD subunit NM_006638 2.71
transducin (beta)-like 1 AAT74134 2.70
cytochrome c oxidase subunit Vb A557312 2.70
RAB5B, member RAS oncogene family AF267863 2.68
RAP2B, member of RAS oncogene family NM_002886 2.67
ATP-binding cassette, sub-family D (ALD), member 3 NM_002858 2.67
Consensus includes gb:AK025724.1 /DEF=Homo sapiens cDNA, FLJ22071 fls, clone HEP11691, /FEA=mRNA /DB_XREF=gi:10438333 /UG=Hs.326248 Homo sapiens cDNA AK025724 2.67
solute carrier family 21 (organic anion transporter), member 6 AB026257 2.65
Consensus includes gb:AW971134 /FEA=EST /DB_XREF=gi:8160979 /DB_XREF=est:EST383221 /UG=Hs.292245 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAM1 AW971134 2.64
orphan seven-transmembrane receptor, chemokine related NM_016557 2.61
Consensus includes gb:AL050065.1 /DEF=Homo sapiens mRNA, cDNA DKFZp566M043 (from clone DKFZp566M043), /FEA=mRNA /DB_XREF=gi:4884295 /UG=Hs.2125 AL050065 2.61
PAX transcription activation domain interacting protein 1 like A1357401 2.60
cytoskeleton-associated protein 4 NM_006825 2.59
KIAA0653 protein, B7-like protein AF289028 2.59
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 6 AF200738 2.59
epithelial membrane protein 3 NM_001425 2.59
ribosomal protein L37a BE85772 2.59
Consensus includes gb:A1345238 /FEA=EST /DB_XREF=gi:4082444 /DB_XREF=est:db1807.x1 /CLONE=IMAGE:2060725 /UG=Hs.111334 ferritin, light polypeptide A1345238 2.59
ATPase, H+ transporting, lysosomal (vacuolar proton pump) 9KD A1252582 2.58
Bloom syndrome NM_000057 2.57
cytochrome P450 isoform 4F12 NM_023944 2.56
H3 histone family, member C NM_003531 2.56
hypothetical protein FLJ22009 NM_024745 2.56
glycophorin E NM_002102 2.55
hypothetical protein FLJ10298 NM_018050 2.55
endomucin-1 NM_016241 2.55
cyclin G2 AW134535 2.55
hexokinase 2 A1761561 2.54
polymerase (DNA directed), eta NM_006502 2.53
culin 5 NM_003478 2.53
PI-3-kinase-related kinase SMG-1 BE000837 2.52
helicase KIAA0054 NM_014104 2.50
M10098 Human 18S rRNA gene, complete (-5'..M..3' represent transcript regions 5 prime, Middle, and 3 prime respectively) NM_014877 2.50
KIAA0869 protein M10098 2.50
hypothetical protein FLJ20897 NM_015377 2.49
gamma-aminobutyric acid (GABA) A receptor, alpha 5 A1335509 2.49
translation initiation factor IF2 BF966183 2.49
SCAN domain-containing 2 BE138647 2.48
inhibin, beta C AF244812 2.47
M10098 Human 18S rRNA sequence, length 1969 bases, 3 prime target bases 1293-1938 NM_005538 2.47
cysteine-rich motor neuron 1 M10098 2.47
mitochondrial ribosomal protein S12 BG546884 2.47
hypothetical protein FLJ10357 R68573 2.47
mamian-binding lectin serine protease 1 (C41C2 activating component of Re-active factor) NM_018071 2.47
thiopurine S-methyltransferase U12387 2.45
sorting nexin 4 AA523345 2.45
golgii associated, gamma adaptin ear containing, ARF binding protein 1 AW001443 2.44
Escherichia coli /REF=J04423 /DEF=E coli bioB gene biotin synthetase corresponding to nucleotides 2772-3004 of J04423 /LEN=1114 (-5'..-M..-3' represent transcript region) J04423 2.44
methionine adenosyltransferase II, alpha AW301881 2.43
Consensus includes gb:AK022473.1 /DEF=Homo sapiens cDNA FLJ12411 fls, clone MAMMA1002964, /FEA=mRNA /DB_XREF=gi:10433882 /UG=Hs.256722 Homo sapiens cDNA AK022473 2.43
eukaryotic translation initiation factor 5A AA393940 2.42
tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains; Huntingtin interacting protein NM_021980 2.42
J04423 E coli bioB gene biotin synthetase (-5'..-M..-3' represent transcript regions 5 prime, Middle, and 3 prime respectively) J04423 2.42
SH3-domain GRB2-like 3 AF036269 2.42

hypothetical protein FLJ13078	AK023140	2.41
serine/threonine-protein kinase PRP4 homolog	AA156948	2.41
harakiri, BCL2 interacting protein (contains only BH3 domain)	U78376	2.41
Consensus includes gb:AK021505.1 /DEF=Homo sapiens cDNA FLJ11443 fls, clone HEMBA1001330 /FEA=mRNA /DB_XREF=gi:10432701 /UG=Hs.297945 Homo sapie	AK021505	2.40
glioma pathogenesis-related protein	U16307	2.40
artemis protein	AK022922	2.40
phosphodiesterase 10A	AF127480	2.38
ubiquitin specific protease 15	NM_006069	2.38
TGF-beta-induced factor (TALE family homeobox)	NM_003244	2.38
PRO0478 protein	NM_014129	2.38
artemis protein	NM_022487	2.37
Consensus includes gb:A015947 /FEA=EST /DB_XREF=gi:5635802 /DB_XREF=ast:w96e01.x1 /CLONE=IMAGE:2379096 /UG=Hs.28212 ESTs	A015947	2.37
triple functional domain (PTPRF interacting)	AL161955	2.36
thromboxane A2 receptor	NM_001060	2.36
KIAA1655 protein	AB051442	2.36
Consensus includes gb:BF942161 /FEA=EST /DB_XREF=gi:12359481 /DB_XREF=ast:nae87g10.x1 /CLONE=IMAGE:4118994 /UG=Hs.302797 ESTs	BF942161	2.35
DnaJ (Hsp40) homolog, subfamily A, member 1	AL534104	2.35
hypothetical protein FLJ22479	NM_024900	2.35
transforming, acidic coiled-coil containing protein 2	AF220152	2.35
KIAA0446 gene product	AB007915	2.34
Consensus includes gb:AL157484.1 /DEF=Homo sapiens mRNA, cDNA DKFZp762M127 (from clone DKFZp762M127) /FEA=mRNA /DB_XREF=gi:7018527 /UG=Hs.224E	N92920	2.33
kinectin 1 (kinesin receptor)	BF589024	2.33
colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)	L28349	2.32
DEADH (Asp-Glu-Ala-Asp/His) box binding protein 1	AL348378	2.32
hypothetical protein FLJ23548	NM_024590	2.32
MAD, mothers against decapentaplegic homolog (Drosophila) interacting protein, receptor activation anchor	NM_007323	2.32
Consensus includes gb:BC004344.1 /DEF=Homo sapiens, clone IMAGE:3633354, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3633354) /DB_XI	BC004344	2.32
Consensus includes gb:R33984 /FEA=EST /DB_XREF=gi:789822 /DB_XREF=ast:y74c03.1 /CLONE=IMAGE:135460 /UG=Hs.286681 Homo sapiens cDNA FLJ11022 fls	R33984	2.31
prostate derived STE20-like kinase PSK	NM_016151	2.30
putative protein O-mannosyltransferase	NM_013382	2.30
Consensus includes gb:AK021440.1 /DEF=Homo sapiens cDNA FLJ11378 fls, clone HEMBA1000456 /FEA=mRNA /DB_XREF=gi:10432625 /UG=Hs.6937 hypothetical pr	AK021440	2.30
Consensus includes gb:AL080180.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434M054 (from clone DKFZp434M054) /FEA=mRNA /DB_XREF=gi:5262622 /UG=Hs.274E	AL080180	2.30
uncharacterized hypothetical protein HT011	BE565675	2.29
S-adenosylmethionine decarboxylase 1	NM_001634	2.29
Consensus includes gb:BE786164 /FEA=EST /DB_XREF=gi:10207362 /DB_XREF=ast:6014273F1 /CLONE=IMAGE:3877146 /UG=Hs.80285 Homo sapiens mRNA, cD	BE786164	2.29
Consensus includes gb:BC005385.1 /DEF=Homo sapiens, clone IMAGE:3829438, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3829438) /DB_XI	BC005385	2.28
Consensus includes gb:AW971254 /FEA=EST /DB_XREF=gi:8161099 /DB_XREF=ast:EST383343 /UG=Hs.178433 ESTs	AW971254	2.28
sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3A	NM_006080	2.28
Consensus includes gb:AF164963.1 /DEF=Homo sapiens tumor antigen NA88-A, pseudogene, complete sequence. /FEA=mRNA /DB_XREF=gi:5901726 /UG=Hs.306576 t	AF164963	2.28
immunoglobulin heavy constant mu	S55735	2.28
actin-related protein 3-beta	NM_020445	2.28
CDC14 cell division cycle 14 homolog A (S. cerevisiae)	NM_003672	2.27
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 17 (72KD)	NM_030881	2.27
PABP-interacting protein 2	AL043487	2.26
DnaJ (Hsp40) homolog, subfamily B, member 9	AL080081	2.26
alcohol dehydrogenase 1C (class I), gamma polypeptide	NM_006069	2.25
ectotropic viral integration site 2A	NM_014210	2.25
Consensus includes gb:A126492 /FEA=EST /DB_XREF=gi:3595006 /DB_XREF=ast:q82n06.x1 /CLONE=IMAGE:1736027 /UG=Hs.104258 Homo sapiens mRNA, exon 1	A126492	2.24
N-myristoyltransferase 2	J04423	2.24
J04423 E coli bioB gene biotin synthetase (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	J04423	2.24
phosducin-like	NM_005388	2.24
hypothetical protein FLJ22558	NM_022747	2.24
peptidyl-prolyl isomerase G (cyclophilin G)	AW340788	2.23
KIAA0469 gene product	NM_014851	2.23
putative lymphocyte G0/G1 switch gene	NM_015714	2.23
Consensus includes gb:BE830512 /FEA=EST /DB_XREF=gi:10456588 /DB_XREF=ast:RC6-GN0071-160800-021-D01 /UG=Hs.168732 ESTs	BE830512	2.23
A kinase (PRKA) anchor protein 1	BC000729	2.23
aldohydroxy dehydrogenase 4 family, member A1	U24287	2.23
Consensus includes gb:BF573849 /FEA=EST /DB_XREF=gi:11647561 /DB_XREF=ast:802132053F1 /CLONE=IMAGE:4271340 /UG=Hs.98343 ESTs, Weakly similar to AL	BF573849	2.22
ATPase, Class I, type BB, member 1	BC025686	2.22
Consensus includes gb:A701158 /FEA=EST /DB_XREF=gi:4989056 /DB_XREF=ast:we10f09.x1 /CLONE=IMAGE:2340713 /UG=Hs.6550 Homo sapiens cDNA: FLJ23227	A701158	2.22
stress 70 protein chaperone, microsome-associated, 60kD	A1718418	2.22

cytochrome P450, subfamily 11C (mephenytoin 4-hydroxylase), polypeptide 19	X65962	2.08
choline/ethanolaminephosphotransferase	NM_006090	2.07
KIAA0304 gene product	AF105279	2.07
endase 1, (alpha)	U88968	2.07
H2A histone family, member X	H51429	2.07
ALEX3 protein	NM_016607	2.07
Consensus includes gb:AK026825.1 /DEF=Homo sapiens cDNA: FLJ23172 fis, clone LNG10005. /FEA=mRNA /DB_XREF=gi:10439771 /UG=Hs.306885 Homo sapiens cDNA	AK026825	2.07
hypothetical protein FLJ20059	NM_017644	2.07
potassium inwardly-rectifying channel, subfamily J, member 8	NM_004982	2.07
early lymphoid activation protein	L22650	2.07
KIAA0874 protein	X80821	2.06
somatostatin receptor 4	NM_001052	2.06
lankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase 2	NM_025235	2.06
immunoglobulin heavy constant gamma 3 (G3m marker)	BF002659	2.06
transmembrane activator and CAML interactor	NM_012452	2.05
RAO54, S. cerevisiae, homolog of B	NM_012415	2.05
polymerase (RNA) II (DNA directed) polypeptide B (140kD)	AW770896	2.05
FK506 binding protein 8 (38kD)	N85418	2.05
translational initiation factor IF2	BG281322	2.05
Consensus includes gb:AL050154.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586L0120 (from clone DKFZp586L0120) /FEA=mRNA /DB_XREF=gi:4884366 /UG=Hs.985 AL050154	AL050154	2.05
Consensus includes gb:AU147017 /FEA=EST /DB_XREF=gi:11008538 /DB_XREF=est:AU147017 /CLONE=HEM8B1002152 /UG=Hs.301905 Homo sapiens cDNA FLJ141	AL147017	2.05
G protein-coupled receptor 1	AL046992	2.05
protease, serine, 21 (testisin)	NM_006799	2.05
hypothetical protein FLJ10496	NM_018114	2.05
pregnancy specific beta-1-glycoprotein 11	NM_002785	2.05
RAN binding protein 1	AI221318	2.04
hypothetical protein FLJ12151	AK022213	2.04
Consensus includes gb:AK022174.1 /DEF=Homo sapiens cDNA FLJ12112 fis, clone MAMMA1000043. /FEA=mRNA /DB_XREF=gi:10433510 /UG=Hs.124126 Homo sapiens clone 24438	AK022174	2.04
transcription factor 4	AK026674	2.04
SRY (sex determining region Y-box 4)	AI899477	2.04
Consensus includes gb:AC004460 /DEF=Homo sapiens PAC clone RPS-1086D14 /FEA=CDS /DB_XREF=gi:2981263 /UG=Hs.307352 Homo sapiens PAC clone RPS-1086	AC004460	2.03
Consensus includes gb:AK023918.1 /DEF=Homo sapiens cDNA FLJ13856 fis, clone THYRO1000988. /FEA=mRNA /DB_XREF=gi:10436003 /UG=Hs.288489 Homo sapiens	AK023918	2.03
SEC24 related gene family, member D (S. cerevisiae)	NM_014822	2.03
hypothetical protein FLJ11338	NM_018393	2.03
checkpoint suppressor 1	AA860806	2.03
Consensus includes gb:AF070647.1 /DEF=Homo sapiens clone 24438 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3283921 /UG=Hs.124126 Homo sapiens clone 24438	AF070647	2.03
translin	AI659160	2.02
neurophilin 2	AA295257	2.02
crystallin-rich motor neuron 1	NM_018441	2.02
S100 calcium binding protein A11 (calgizarin)	NM_005620	2.02
gb:NM_017648.1 /DEF=Homo sapiens hypothetical protein FLJ20063 (FLJ20063), mRNA. /FEA=mRNA /GEN=FLJ20063 /PROD=hypothetical protein FLJ20063 /DB_XREF=	NM_017648	2.02
PC4 and SFRS1 interacting protein 1	AF068482	2.02
immunoglobulin heavy constant mu	S74639	2.02
Consensus includes gb:AW301808 /FEA=EST /DB_XREF=gi:5711483 /DB_XREF=est:rs5611.x1 /CLONE=IMAGE2764172 /UG=Hs.150551 ESTs, Weakly similar to ALU	AW301808	2.02
ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome)	BF588511	2.01
adectin 2 (beta)	NM_017482	2.01
thyroid hormone receptor-associated protein, 95-kD subunit	BG338608	2.01
HCF-binding transcription factor Zhanfiei	NM_018810	2.01
protein phosphatase 2, regulatory subunit B (B56), gamma isoform	AI206560	2.01
cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)	AW772123	2.01
F-box only protein 21	AK023368	2.00
gb:AF060693.1 /DEF=Homo sapiens clone HQ0117 PRO0117 mRNA, complete cds. /FEA=mRNA /PROD=PRO0117 /DB_XREF=gi:6690166 /UG=Hs.283919 Homo sapiens	AF060693	2.00
hypothetical protein PRO2849	AK001699	2.00
Epstein-Barr virus induced gene 3	NM_022335	2.00
ATPase, Ca++ transporting, plasma membrane 1	NM_003755	2.00
tripartite motif-containing 2	L14561	2.00
brain-specific angiogenesis inhibitor 3	NM_015271	2.00
Consensus includes gb:X78262.1 /DEF=H sapiens mRNA for TRES. /FEA=mRNA /DB_XREF=gi:587440 /UG=Hs.302178 H.sapiens mRNA for TRES	AB011122	2.00
neuregulin 2	X78262	2.00
v-myc myelocytomatosis viral oncogene homolog 2 (avian)	NM_013984	2.00
calpain 9 (nCL-4)	NM_005377	1.99
zinc finger protein 281	AB038463	1.99
	NM_012482	1.99

hypothetical protein LOC57187
 hypothetical protein FLJ13166
 gastric inhibitory polypeptide receptor
 activation-induced cytidine deaminase
 caspase 8, apoptosis-related cysteine protease
 hypothetical protein FLJ20837
 dickkopf homolog 2 (Xenopus laevis)
 lipopolysaccharide specific response-68 protein
 protein phosphatase, EF hand calcium-binding domain 2
 hypothetical protein FLJ14348
 membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE, high affinity 1, receptor for beta polypeptide)
 Escherichia coli (REF=J04423) DEF=EE coil b10c protein corresponding to nucleotides 4609-4883 of J04423 (LEN=777 (-5 and -3 represent transcript regions 5 prime and 3 J04423
 Consensus includes gb:BF035278 /FEA=EST /DB_XREF=gi:10743008 /DB_XREF=est:601457185F1 /CLONE=IMAGE:3860633 /UG=Hs.20325 ESTs. Moderately similar to
 KIAA1102 protein
 mannosyl (alpha-1,6)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
 ornithine decarboxylase anilzyme inhibitor
 BLU protein
 phosphatidylinositol-4-phosphate 5-kinase, type II, alpha
 clone FLB3816
 thyroid hormone receptor interactor 11
 small inducible cytokine subfamily A (Cys-Cys), member 18, pulmonary and activation-regulated
 hypothetical protein FLJ13162
 Homer, neuronal immediate early gene, 1B
 hypothetical protein FLJ14310
 Consensus includes gb:U43804.1 /DEF=Human unidentified mRNA, partial sequence. /FEA=mRNA /DB_XREF=gi:1171236 /UG=Hs.159801 Human unidentified mRNA, pt U43804
 hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 2
 lipin 2
 Consensus includes gb:AL050122.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586E121 (from clone OKFZp586E121) /FEA=mRNA /DB_XREF=gi:4884330 /UG=Hs.2745 AL050122
 gb:U96291.1 /DEF=Homo sapiens Ig kappa light chain variable region (VKII-A23) mRNA, complete cds. /FEA=mRNA /GEN=VKII-A23 /PROD=ig kappa light chain variable 1 U96291
 P3ECSL
 patched homolog (Drosophila)
 WW domain binding protein 4 (fornin binding protein 21)
 neural cell adhesion molecule 2
 hypothetical protein FLJ10254
 ubiquitin-conjugating enzyme E2i (UBCS homolog, yeast)
 peptide YY, 2 (seminal plasma)
 DnaJ (Hsp40) homolog, subfamily C, member 8
 KIAA0692 protein
 UDP-glucuronic acid/UDP-N-acetylglucosamine dual transporter
 hypothetical protein FLJ20084
 beta-amyloid binding protein precursor
 glucocorticoid modulatory element binding protein 2
 coagulation factor V (proaccelerin, labile factor)
 KIAA0256 gene product
 Consensus includes gb:AK000854.1 /DEF=Homo sapiens cDNA FLJ10002 /fs, clone HEMBA1000046. /FEA=mRNA /DB_XREF=gi:7021188 /UG=Hs.236522 Homo sapien
 mitochondrial ribosomal protein L44
 Consensus includes gb:AK022219.1 /DEF=Homo sapiens cDNA FLJ12157 /fs, clone MAMMA1000500. /FEA=mRNA /DB_XREF=gi:10433569 /UG=Hs.98812 Homo sapier
 jumortil homolog (mouse)
 homeo box C5
 cAMP responsive element binding protein-like 1
 Alu-binding protein with zinc finger domain
 KIAA0318 gene product
 golgi autoantigen, golgin subfamily a, 4
 hypothetical protein dJ482023.2
 solute carrier family 16 (monocarboxylic acid transporters), member 4
 tripartite motif-containing 5
 integral membrane protein 2B
 ATPase, Cu++ transporting, alpha polypeptide (Menkes syndrome)
 cyclin E2
 EphB3
 hypothetical protein FLJ20097
 DnaJ (Hsp40) homolog, subfamily B, member 9
 polymerase (RNA) III (DNA directed) (32kD)
 NM_006467

aldehyde dehydrogenase 1 family, member B1
 Consensus includes gb:AA017721 /FEA=EST /DB_XREF=gi:1479910 /DB_XREF=est:z639f11.31 /CLONE=IMAGE:361389 /UG=Hs.49.117 Homo sapiens mRNA, cDNA
 nuclear transcription factor Y, alpha
 calcitonin receptor
 tubby like protein 3
 hypothetical protein
 hypothetical protein FLJ11328
 golgi associated, gamma adaptin ear containing, ARF binding protein 2
 Consensus includes gb:AK023783.1 /DEF=Homo sapiens cDNA FLJ13721 fls, done PLACE2000450, /FEA=mRNA /DB_XREF=gi:10435820 /UG=Hs.289035 Homo sapiens
 succinate dehydrogenase complex, subunit C, integral membrane protein, 15KD
 hypothetical protein FLJ20019
 KIAA1110 protein
 Consensus includes gb:BC049571 /FEA=EST /DB_XREF=gi:13457287 /DB_XREF=est:602540396F1 /CLONE=IMAGE:4671599 /UG=Hs.289068 Homo sapiens cDNA FLJ
 zinc finger protein 43 (H1F6)
 Fc fragment of IgG, low affinity IIb, receptor for (CD16)
 hypothetical protein similar to small G proteins, especially RAP-2A
 synovial sarcoma translocation gene on chromosome 18-like 1
 Consensus includes gb:AW576457 /FEA=EST /DB_XREF=est:U1-HF-BR0p-slj-9-01-Q-U1.s1 /CLONE=IMAGE:3074658 /UG=Hs.20952 Homo sapiens
 prostate differentiation factor
 KIAA1827 protein
 trichothiodystrophy syndrome 1
 dynamin 2
 dopamine receptor D4
 Consensus includes gb:AK025095.1 /DEF=Homo sapiens cDNA: FLJ21442 fls, done COL04429, highly similar to HSA237839 Homo sapiens mRNA for hypothetical protein
 NG37 protein
 protein kinase, cAMP-dependent, regulatory, type II, alpha
 hypothetical protein DKFZp34D0421
 solute carrier family 9 (sodium/hydrogen exchanger), isoform 2
 phospholamban
 Consensus includes gb:AV684859 /FEA=EST /DB_XREF=gi:10286722 /DB_XREF=est:AV684859 /CLONE=GKCDJAO3 /UG=Hs.251334 ESTs, Moderately similar to ALU6
 glutamate receptor, ionotropic, kainate 4
 male germ cell-associated kinase
 fatty-acyl-Coenzyme A ligase, long-chain 3
 Fc fragment of IgE, high affinity I, receptor for: alpha polypeptide
 neuronal Shc
 hypothetical protein FLJ23412
 transmembrane protein 5
 sorting nexin 15
 G-2 and S-phase expressed 1
 Consensus includes gb:AW970584 /FEA=EST /DB_XREF=gi:8160429 /DB_XREF=est:EST382685 /UG=Hs.291033 ESTs
 KIAA0480 gene product
 Consensus includes gb:AW451711 /FEA=EST /DB_XREF=gi:6992487 /DB_XREF=est:U1-H-B13-ale-c-02-Q-U1.s1 /CLONE=IMAGE:2736386 /UG=Hs.313760 ESTs
 protein tyrosine kinase 9
 monoamine oxidase A
 homeo box A10
 nuclear factor I/B
 KIAA0415 gene product
 GABA(A) receptor-associated protein like 1
 hypothetical protein PRQ0082
 bone morphogenetic protein receptor, type IB
 Consensus includes gb:A732770 /FEA=EST /DB_XREF=gi:5053383 /DB_XREF=est:z78d05.x5 /CLONE=IMAGE:809865 /UG=Hs.328688 ESTs, Moderately similar to AL
 pleomorphic adenoma gene 1
 KIAA0341 protein
 Consensus includes gb:AL312703.1 /DEF=Homo sapiens mRNA; cDNA DKFZp667H046 (from clone DKFZp667H046) /FEA=mRNA /GEN=DKFZp667H046 /PROD=hypoth
 Consensus includes gb:A734156 /FEA=EST /DB_XREF=gi:5055269 /DB_XREF=est:mc79004.y5 /CLONE=IMAGE:783559 /UG=Hs.172895 ESTs
 tripeptidyl peptidase II
 glucan (1,4-alpha-), branching enzyme 1 (glycogen branching enzyme, Andersen disease, glycogen storage disease type IV)
 intracellular hyaluronan-binding protein
 LIM domain protein
 mitogen-activated protein kinase 6
 SRY (sex determining region Y)-box 21
 hypothetical protein SP192
 Wilms tumor 1

laminin, beta 1
 Consensus includes gb:A002715 /FEA=EST /DB_XREF=gi:3203129 /DB_XREF=est:an20d12.s1 /CLONE=IMAGE:1699223 /UG=Hs.130315 ESTs
 general transcription factor IIF, polypeptide 3 (34kD subunit)
 putative zinc finger protein from EUROMAGE 566589
 killer cell lectin-like receptor subfamily D, member 1
 doublesex and mab-3 related transcription factor 1
 estrogen receptor 2 (ER beta)
 Consensus includes gb:AW301806 /FEA=EST /DB_XREF=gi:6711463 /DB_XREF=est:zx56e11.x1 /CLONE=IMAGE:2764172 /UG=Hs.150551 ESTs, Weakly similar to ALU AW301806
 similar to rat nuclear ubiquitin casein kinase 2
 antigen identified by monoclonal antibody MRC OX-2
 glycoprotein IX (platelet)
 B-cell CLL/lymphoma 7A
 basic helix-loop-helix domain containing, class B, 3
 IMP (inosine monophosphate) dehydrogenase 1
 PC3-96 protein
 cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4)
 matrix metalloproteinase 25
 death-associated protein kinase 2
 Consensus includes gb:AF222691.1 /DEF=Homo sapiens Alu repeat (LNX1) mRNA sequence. /FEA=mRNA /DB_XREF=gi:12655850 /UG=Hs.307008 Homo sapiens Alu r. AF222691
 RAB3A, member RAS oncogene family
 CED-8 protein
 Consensus includes gb:AK021609.1 /DEF=Homo sapiens cDNA FLJ11547 fls, clone HEMBA1002934. /FEA=mRNA /DB_XREF=gi:10432824 /UG=Hs.306605 Homo sapiens
 ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome
 nephrosis 1, congenital, Finnish type (nephtrin)
 HSPC039 protein
 BRCA1 associated protein
 3-phosphoinositide dependent protein kinase-1
 Consensus includes gb:BE672313 /FEA=EST /DB_XREF=gi:10032854 /DB_XREF=est:7a59b10.x1 /CLONE=IMAGE:3223003 /UG=Hs.34054 Homo sapiens cDNA: FLJ22
 Consensus includes gb:AL136105 /DEF=Human DNA sequence from clone RP4-670F13 on chromosome 1q42.2-43. Contains an enolase 1, (alpha) (ENO1) pseudogene, AL117387
 PDZ domain containing guanine nucleotide exchange factor (GEF) 1
 hypothetical protein FLJ23142
 a disintegrin-like and metalloprotease (repolysin type) with thrombospondin type 1 motif, 1
 KIAA1045 protein
 craniofacial development protein 1
 G protein-coupled receptor 37 (endothelin receptor type B-like)
 CS box-containing WD protein
 KIAA1685 protein
 hypothetical protein FLJ22748
 J04423 E. coli bioC protein (-5 and -3 represent transcript 1 regions 5 prime and 3 prime respectively)
 signal transduction protein (SH3 containing)
 KIAA0368 protein
 hypothetical protein FLJ13448
 melanocortin 1 receptor (alpha melanocyte stimulating hormone receptor)
 Consensus includes gb:U63542.1 /DEF=Human putative FAP protein mRNA, partial cds. /FEA=mRNA /PROD=FAP protein /DB_XREF=gi:1890646 /UG=Hs.166101 Human
 likely ortholog of mouse polydrom
 heterogeneous nuclear ribonucleoprotein H3 (2H8)
 kangal 1 (suppression of tumorigenicity 6, prostate, COB2 antigen (R2 leukocyte antigen, antigen detected by monoclonal and antibody 1A4))
 carboxypeptidase B1 (tissue)
 acute carrier family 13 (sodium/sulfate symporters), member 1
 hypothetical protein FLJ12873
 J04423 E. coli bioD gene dehydrobiolin synthetase (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)
 KIAA0594 protein
 high-mobility group (nonhistone chromosomal) protein 4
 neuronal thread protein
 GK001 protein
 heterogeneous nuclear ribonucleoprotein H1 (H)
 glutaminase
 Consensus includes gb:AK026484.1 /DEF=Homo sapiens cDNA: FLJ22831 fls, clone KIAA1461. /FEA=mRNA /DB_XREF=gi:10439356 /UG=Hs.321666 Homo sapiens cD
 pre-mRNA processing factor 31 homolog (yeast)
 mitochondrial ribosomal protein L18
 macrophage stimulating, pseudogene 9
 dopamine receptor D2

double homeobox, 4	D38024	1.85
odd-skipped-related 2A protein	A1811238	1.85
zinc finger protein 42 (myeloid-specific retinoic acid- responsive)	M58297	1.85
Interleukin 8 receptor, beta	NM_001557	1.85
G protein-coupled receptor 4	NM_005282	1.85
KIAA0007 protein	D26488	1.85
caspase 5, apoptosis-related cysteine protease	NM_004347	1.85
rab3 GTPase-activating protein, non-catalytic subunit (150KD)	AK021928	1.85
carboxypeptidase M	NM_001874	1.85
alkaline phosphatase, intestinal	NM_001631	1.84
myosin, heavy polypeptide 13, skeletal muscle	NM_003802	1.84
hypothetical protein PRO2533	NM_018629	1.84
decay accelerating factor for complement (CD55, Cromer blood group system)	BC001288	1.84
hypothetical protein PRO2800	NM_018635	1.84
glycophorin A (includes MN blood group)	U00170	1.84
Consensus includes gb:AK024136.1 /DEF=Homo sapiens cDNA FLJ14074 fs, clone HEMBB1001869, /FEA=mRNA /DB_XREF=gi:10436442 /UG=Hs.141208 Homo sapie	AK024136	1.84
hypothetical protein MGC3196	NM_024084	1.84
neuropeptide Y receptor Y6 (pseudogene)	U59431	1.84
Lsm1 protein	NM_014462	1.84
integrin, beta 5	AU35208	1.84
hypothetical protein FLJ23447	NM_024825	1.84
dipeptidase 1 (renal)	NM_004413	1.84
Consensus includes gb:AK025038.1 /DEF=Homo sapiens cDNA: FLJ21385 fs, clone COL03356, /FEA=mRNA /DB_XREF=gi:10437474 /UG=Hs.306750 Homo sapiens cC AK025038	AK025038	1.84
Consensus includes gb:A1570199 /FEA=EST /DB_XREF=gi:4533573 /DB_XREF=est:to76b07.x1 /CLONE=IMAGE:2184181 /UG=Hs.182183 Homo sapiens mRNA for cald, A1570199	AK025171	1.84
gb:BC000108.1 /DEF=Homo sapiens, Similar to Nedd-4-like ubiquitin-protein ligase, clone MGC:2079, mRNA, complete cds. /FEA=mRNA /PROD=Similar to Nedd-4-like u	BC000108	1.84
EphB4	BC004264	1.84
translation initiation factor IF2	BE138647	1.84
KIAA0138 gene product	D50928	1.83
immunoglobulin superfamily, member 4	AL519710	1.83
connective tissue growth factor	M92934	1.83
sex comb on midleg-like 2 (Drosophila)	NM_006089	1.83
Consensus includes gb:AK021571.1 /DEF=Homo sapiens cDNA FLJ11509 fs, clone HEMBA1002166, /FEA=mRNA /DB_XREF=gi:10432777 /UG=Hs.285093 Homo sapie	AK021571	1.83
IGF-II mRNA-binding protein 2	NM_006548	1.83
Consensus includes gb:AK027193.1 /DEF=Homo sapiens cDNA: FLJ23540 fs, clone LNG08239, /FEA=mRNA /DB_XREF=gi:10440282 /UG=Hs.293331 Homo sapiens cC AK027193	AK027193	1.83
eukaryotic translation initiation factor 4E	NM_001968	1.83
taste receptor, type 2, member 16	NM_016945	1.83
hypothetical protein FLJ10687	NM_018178	1.83
Consensus includes gb:AW301937 /FEA=EST /DB_XREF=gi:8711614 /DB_XREF=est:tr85b03.x1 /CLONE=IMAGE:2766893 /UG=Hs.138038 ESTs	AW301937	1.83
Consensus includes gb:BE786598 /FEA=EST /DB_XREF=gi:10207796 /DB_XREF=est:601478824F1 /CLONE=IMAGE:3881717 /UG=Hs.101474 KIAA0807 protein	BE786598	1.83
phospholipid scramblase 2	NM_020359	1.83
N-myc downstream-regulated gene 2	AU358939	1.83
SPARC-like 1 (mest8, hev1n)	NM_004684	1.83
RAB36, member RAS oncogene family	AF133588	1.83
Consensus includes gb:AK024103.1 /DEF=Homo sapiens cDNA FLJ14046 fs, clone HEMBA1006461, /FEA=mRNA /DB_XREF=gi:10436406 /UG=Hs.142677 Homo sapie	AK024108	1.83
TXK tyrosine kinase	AW276646	1.83
mitogen-activated protein kinase 8 interacting protein 2	NM_016431	1.83
RNA-binding protein gene with multiple splicing	DB4109	1.82
trinucleotide repeat containing 15	AB014542	1.82
similar to rab11-binding protein	NM_019045	1.82
complement component 7	NM_000587	1.82
keratin 18	AL031903	1.82
hypothetical protein FLJ21625	NM_025039	1.82
pleckstrin homology-like domain, family A, member 1	A1795908	1.82
Consensus includes gb:AW973253 /FEA=EST /DB_XREF=gi:8163100 /DB_XREF=est:EST385351 /UG=Hs.292689 ESTs	AW973253	1.82
novel putative protein similar to YIL091C yeast hypothetical 84 kD protein from SGA1-KTR7	A1770084	1.82
hypothetical protein FLJ10486	BF591163	1.82
short stature homeobox 2	NM_006884	1.82
insulin	NM_000207	1.82
atractin	AL132773	1.81
retinoblastoma binding protein 6	NM_006910	1.81
Consensus includes gb:AL049260.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564E233 (from clone DKFZp564E233), /FEA=mRNA /DB_XREF=gi:4500007 /UG=Hs.3020 /AL049260	AL049260	1.81
Klotho	NM_004795	1.81
glycoprotein M6B	A1479030	1.81

Cdc42 guanine exchange factor (GEF) 9	1.81
DKFZ564J102 protein	1.81
RAP2B, member of RAS oncogene family	1.81
CASP8 and FADD-like apoptosis regulator	1.81
Intermediate filament protein syncollin	1.81
hypothetical protein FLJ20048	1.81
protein phosphatase 2 (lomerly 2A), regulatory subunit B' (PR 72), alpha isoform and (PR 130), beta isoform	1.81
MDS024 protein	1.81
unc-51-like Kinase 2 (C. elegans)	1.81
DEAD/HD (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)	1.81
hormo box C8	1.81
calcium/calmodulin-dependent serine protein kinase (IMAGUK family)	1.81
myosin, heavy polypeptide 1, skeletal muscle, adult	1.81
HMT1 hnRNP methyltransferase-like 1 (S. cerevisiae)	1.81
transducer of ERBB2, 2	1.81
lysine aminotransferase	1.81
sirolimus (GdL115, leucosialin, CD43)	1.81
Consensus includes gb:AL050136.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586L141 (from done DKFZp586L141). /FEA=mRNA /DB_XREF=gi:4884346 /UG=Hs. 14094	1.80
sialic acid binding Ig-like lectin 8	1.80
tumor protein p53 (Li-Fraumeni syndrome)	1.80
oxidative stress induced like	1.80
Consensus includes gb:BG290532 /FEA=EST /DB_XREF=gi:13047560 /DB_XREF=est:602398395F1 /CLONE=IMAGE:4517206 /UG=Hs. 11210 ESTs, Moderately similar t	1.80
bromodomain, testis-specific	1.80
prostatic binding protein	1.80
alpha-2-glycoprotein 1, zinc	1.80
erythrocyte membrane protein band 7.2 (stomatin)	1.80
coagulation factor II (thrombin) receptor	1.80
Consensus includes gb:AK026913.1 /DEF=Homo sapiens cDNA: FLJ23260 fs, clone COL05804, highly similar to HSU9011 Human clone 23652 mRNA sequence. /FEA=	1.80
CLL17 protein	1.80
neurotrophic tyrosine kinase, receptor, type 1	1.80
serine protease inhibitor-like, with Kunitz and WAP domains 1 (epnin)	1.80
L39424.B subtilis daps, jpf, jpfG genes corresponding to nucleotides 1358-3197 of L38424 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respective)	1.80
zinc finger protein 161	1.80
homolog of mouse BMP-2 inducible kinase	1.80
CGI-142	1.80
Consensus includes gb:AJ042030 /FEA=EST /DB_XREF=gi:3281224 /DB_XREF=est:ox60a09.x1 /CLONE=IMAGE:1660696 /UG=Hs. 132153 ESTs	1.80
U6 snRNA-associated Sm-like protein	1.80
fatty acid desaturase 2	1.80
actin related protein 2/3 complex, subunit 1B (41 kD)	1.80
serma domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C	1.80
Consensus includes gb:BE326381 /FEA=EST /DB_XREF=gi:9200157 /DB_XREF=est:hw02e08.x1 /CLONE=IMAGE:3181766 /UG=Hs. 18724 Homo sapiens mRNA: cDNA	1.79
chromosome 21 open reading frame 82	1.79
nudin (nucleoside diphosphate linked moiety X)-type motif 7	1.79
Consensus includes gb:AL390026 /DEF=Human DNA sequence from clone RP3-336K20 on chromosome 6 Contains parts of 2 genes for novel proteins, ESTs, STs and	1.79
beta-amyloid binding protein precursor	1.79
peter pan homolog (Drosophila)	1.79
SMART/HDAC1 associated repressor protein	1.79
transmembrane gamma-carboxyglutamic acid protein 4	1.79
interleukin 4	1.79
bromodomain adjacent to zinc finger domain, 1A	1.79
hypothetical protein FLJ13769	1.79
hypothetical protein FLJ10978	1.79
Consensus includes gb:AK023017.1 /DEF=Homo sapiens cDNA FLJ12955 fs, clone NT2RP2005496, moderately similar to ZINC FINGER PROTEIN 135. /FEA=mRNA /DB_XREF=gi:9513716 /DB_XREF=est:hz76b04.x1 /CLONE=IMAGE:3213871 /UG=Hs. 25075 homolog of yeast long chain	1.79
v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian)	1.79
GTP binding protein 2	1.79
mitogen-activated protein kinase 5	1.79
Consensus includes gb:BE467941 /FEA=EST /DB_XREF=gi:9513716 /DB_XREF=est:hz76b04.x1 /CLONE=IMAGE:3213871 /UG=Hs. 25075 homolog of yeast long chain	1.79
DKFZ564B0769 protein	1.79
KIA04186 gene product	1.79
makorin, ring finger protein, 4	1.79
testis specific ankyrin-like protein 1	1.79
transcription elongation factor A (SII), 2	1.79

tyrosine kinase with immunoglobulin and epidermal growth factor homology domains
Williams-Beuren syndrome chromosome region 23
ledlinin 1
SCAN domain-containing 2
v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog
T-box 2
N-myc downstream-regulated gene 2
p21 (CDKN1A)-activated kinase 2
HIC1-related gene on chromosome 22
hypothetical protein FLJ23516
ubiquitin-conjugating enzyme E2 variant 2
apolipoprotein F
Consensus includes gb:AW025578 /FEA=EST /DB_XREF=gi:5879109 /DB_XREF=est:u75b04.x1 /CLONE=IMAGE:2525839 /UG=Hs.100261 Homo sapiens mRNA; cDNA, AW025578
azurocidin 1 (cationic antimicrobial protein 37)
adenomatous polyposis coli
Consensus includes gb:AA573805 /FEA=EST /DB_XREF=gi:2348320 /DB_XREF=est:u75b04.x1 /CLONE=IMAGE:1012888 /UG=Hs.152939 Homo sapiens clone 24630 r
interleukin 18 (interferon-gamma-inducing factor)
hypothetical protein FLJ10701
hypothetical protein FLJ23392
CGI-148 protein
matrix metalloproteinase 10 (stromelysin 2)
DKFZP564A122 protein
CD47 antigen (Rb-related antigen, integrin-associated signal transducer)
Imprinted in Prader-Willi syndrome
Sm protein F
retinoblastoma-like 2 (p130)
signal sequence receptor, alpha (translocin-associated protein alpha)
5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-coupled)
myeloidlymphoid or mixed-lineage leukemia (t(9;22)(q34;q11)) translocated to, 10
hypothetical protein MGC2656
PTPRF interacting protein, binding protein 1 (liprin beta 1)
SmcX homolog, X chromosome (mouse)
interferon regulatory factor 2
KIAA0472 protein
phosphoenolpyruvate carboxykinase 1 (soluble)
DEAD/HA (Asp-Glu-Ala-Asp/His) box polypeptide 34
cyclic fibrosis transmembrane conductance regulator, ATP-binding cassette (sub-family C, member 7)
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily A, member 1
gb:NM_008273.2 /DEF=Homo sapiens small inducible cytokine A7 (monocyte chemoattractant protein 3) (SCYA7), mRNA, /FEA=mRNA /GEN=SCYA7 /PROD=monocyte chem
hypothetical protein PRO2796
Consensus includes gb:NM48832 /FEA=EST /DB_XREF=gi:1188018 /DB_XREF=est:y73d05.s1 /CLONE=IMAGE:279177 /UG=Hs.285625 Homo sapiens mRNA; cDNA DK
hypothetical protein FLJ14166
chymotrypsin C (caldecrin)
heat shock 70kD protein 8
receptor tyrosine kinase-like orphan receptor 1
acetylcholinesterase (YT blood group)
chromosome 6 open reading frame 32
mitogen-activated protein kinase kinase kinase 5
DKFZP566K023 protein
Consensus includes gb:AV753204 /FEA=EST /DB_XREF=gi:10911052 /DB_XREF=est:AV753204 /CLONE=NPD0B08 /UG=Hs.170287 ESTs
hypothetical protein FLJ2849 similar to signal peptidase SPC2223
KIAA1644 protein
T cell activation, increased late expression
KRA8 zinc finger protein KR18
gb:NM_024305.1 /DEF=Homo sapiens hypothetical protein MGC4278 (MGC4278), mRNA, /FEA=mRNA /GEN=MGC4278 /PROD=hypothetical protein MGC4278 /DB_XREF=gi:10433276 /UG=Hs.273759 Homo sapie
neurofilament 3 (150kD medium)
T-box 5
AT2 receptor-interacting protein 1
Consensus includes gb:AK021967.1 /DEF=Homo sapiens cDNA FLJ11905 f1s, clone HEMBB1000050, /FEA=mRNA /DB_XREF=gi:10433276 /UG=Hs.273759 Homo sapie
retinoid X receptor, alpha
cholesterol 25-hydroxylase
UDP-glucose dehydrogenase
KIAA0871 protein

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gem (nuclear organelle) associated protein 4	NM_015487	1.73
melanoma antigen recognized by T cells 2	NM_018194	1.73
transcription factor (p38 interacting protein)	AW003334	1.73
CGI-58 protein	BC006637	1.73
gila maturation factor, beta	NM_004124	1.73
serine/threonine kinase 6	NM_003158	1.73
JM27 protein	NM_007003	1.73
KIAA0874 protein	X80821	1.73
Consensus includes gb:AF339807.1 /DEF=Homo sapiens clone IMAGE:26186, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507345 /UG=Hs.137567 Homo sapiens cDNA AF339807	AF339807	1.73
D site of albumin promoter (albumin D-box) binding protein	D28468	1.73
lymphoid nuclear protein related to AF4	NM_002285	1.73
erythropoietin receptor	M34986	1.73
G protein-coupled receptor 44	NM_004778	1.73
tripartite motif-containing 37	AK022701	1.73
eukaryotic translation initiation factor 4 gamma, 1	BE966878	1.73
cyclin-dependent kinase 6	A1738463	1.73
Consensus includes gb:BG109855 /FEA=EST /DB_XREF=gi:12603361 /DB_XREF=est:50227952.1 /CLONE=IMAGE:4367308 /UG=Hs.49476 Homo sapiens clone TUAJ BG109855	AF035300	1.73
cadherin-like 22	AW173720	1.73
Consensus includes gb:AW173720 /FEA=EST /DB_XREF=gi:6439668 /DB_XREF=est:x1105.x1 /CLONE=IMAGE:2656929 /UG=Hs.271356 ESTs	AW173720	1.73
Consensus includes gb:AL121900 /DEF=Human DNA sequence from clone RPT11-379J5 on chromosome 20 Contains the last exon of the SEC23B gene for Sec23 (S. cere	AL121900	1.73
spectrin, alpha, erythrocytic 1 (elliptocytosis 2)	W87601	1.72
small nuclear ribonucleoprotein polypeptide E	AF131850	1.72
etoposide-induced mRNA	AF131850	1.72
Consensus includes gb:BF693921 /FEA=EST /DB_XREF=gi:11979329 /DB_XREF=est:502082488.F1 /CLONE=IMAGE:4246917 /UG=Hs.180513 ATP-binding cassette, sub	BF693921	1.72
Src-like adaptor	U44403	1.72
5' nucleotidase, mitochondrial	NM_020201	1.72
hypothetical protein FLJ10430	NM_018092	1.72
Consensus includes gb:AL137590.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434K0810 (from clone DKFZp434K0610). /FEA=mRNA /DB_XREF=gi:5808330 /UG=Hs.30	AL137590	1.72
PER011 family member in MHC class I region	NM_005844	1.72
GRO1 oncogene (melanoma growth stimulating activity, alpha)	NM_001511	1.72
Consensus includes gb:W86781 /FEA=EST /DB_XREF=gi:1400509 /DB_XREF=est:zh6a03.s1 /CLONE=IMAGE:416812 /UG=Hs.293736 ESTs	W86781	1.72
chorionic gonadotropin, beta polypeptide	NM_000737	1.72
CDC5 cell division cycle 5-like (S. pombe)	AB007892	1.72
leucine-rich repeat-containing 5	NM_018103	1.72
ubiquitin-like 4	BE543027	1.72
myeloid lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila), translocated to, 4	AL049698	1.72
hairy homolog (Drosophila)	NM_005524	1.72
zinc finger protein, subfamily 1A, 4 (Eos)	NM_022465	1.72
sodium bicarbonate transporter 4	AF243499	1.72
choroideremia-like (Rab escort protein 2)	NM_001821	1.72
Consensus includes gb:A4119423 /FEA=EST /DB_XREF=gi:4265354 /DB_XREF=est:tt28d09.x1 /CLONE=IMAGE:2097521 /UG=Hs.172780 ESTs	A4119423	1.72
lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase)	AW084510	1.72
Consensus includes gb:A1866582 /FEA=EST /DB_XREF=gi:4897876 /DB_XREF=est:td9e05.x1 /CLONE=IMAGE:2268608 /UG=Hs.283824 Homo sapiens Skn-1a1-like m	A1866582	1.72
Consensus includes gb:AW268081 /FEA=EST /DB_XREF=gi:5655111 /DB_XREF=est:tr92e03.x1 /CLONE=IMAGE:2767612 /UG=Hs.193408 ESTs, Weakly similar to ALU	AW268081	1.72
one cut domain, family member 1	U96173	1.72
hypothetical protein FLJ13885	NM_025016	1.72
gb:NM_018541.1 /DEF=Homo sapiens hypothetical protein PRO2832 (PRO2832), mRNA. /FEA=mRNA /GEN=PRO2832 /PROD=hypothetical protein PRO2832 /DB_XREF	NM_018541	1.72
KIAA0328 protein	AB002328	1.72
caveolin 2	AA150110	1.72
phospholipase A2, group VII (platelet-activating factor acetylhydrolase, plasma)	NM_005084	1.72
apolipoprotein C1	W76394	1.72
pleckstrin 2 (mouse) homolog	NM_016445	1.72
hypothetical protein similar to rat vacuolar sorting protein 33a	NM_022916	1.71
cytochrome P450, subfamily X1B (steroid 11-beta-hydroxylase), polypeptide 2	X54741	1.71
complement factor H-related 4	NM_006684	1.71
Consensus includes gb:AF023203.1 /DEF=Homo sapiens homeobox protein Ogl12 (OGL12) mRNA, partial cds. /FEA=mRNA /GEN=OGL12 /PROD=homeobox protein Ogl12	AF023203	1.71
Max-interacting protein	BE502432	1.71
chromosome 5 open reading frame 7	NM_016604	1.71
rat guanine nucleotide dissociation stimulator	A4271559	1.71
melanoma antigen, family D, 2	A1924630	1.71
variable charge, Y chromosome	NM_004879	1.71
H2A histone family, member O	BC001629	1.71
forkhead box G1A	NM_004471	1.71

KIAA1036 protein	NM_014909	1.71
homolog of mouse BMP-2 inducible kinase	AB015331	1.71
CD34 antigen	M61104	1.71
deiodinase, iodothyronine, type I	NM_000792	1.71
ribosome binding protein 1, homolog 180kD (dog)	AA706065	1.71
hypothetical protein FLJ10201	NM_018023	1.71
bridging integrator 2	NM_016293	1.71
mitogen-activated protein kinase kinase 11	NM_002419	1.71
hypothetical protein PRO1496	NM_018603	1.71
integrin, alpha L (antigen CD11A (p180), lymphocyte function-associated antigen 1; alpha polypeptide)	AC002310	1.71
hypothetical protein FLJ10871	NM_018250	1.71
nuclear receptor binding factor-2	AA683074	1.71
retinoic acid repressible protein	AF172066	1.71
decorin	AF138303	1.71
RAD1 homolog (S. pombe)	NM_002853	1.71
glutamate receptor, metabotropic 7	NM_000844	1.71
Consensus includes gb:AF070547.1 /DEF=Homo sapiens clone 24820 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387909 /UG=Hs.146312 Homo sapiens clone 24820 AF070547	AF081459	1.70
origin recognition complex, subunit 5-like (yeast)	AF035594	1.70
Consensus includes gb:AF035594.1 /DEF=Homo sapiens protein kinase C-alpha mRNA, partial 3 UTR. /FEA=mRNA /DB_XREF=gi:3168857 /UG=Hs.279856 Homo sapiens	NM_021163	1.70
hypothetical protein similar to small G proteins, especially RAP-2A	NM_016242	1.70
endomucin-2	NM_001397	1.70
endothelin converting enzyme 1	AF260113	1.70
cytochrome P450 polypeptide 43	NM_014936	1.70
ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function)	AB001328	1.70
solute carrier family 15 (oligopeptide transporter), member 1	NM_008665	1.70
leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 3	AA634446	1.70
Consensus includes gb:AA634446 /FEA=EST /DB_XREF=gi:2557680 /DB_XREF=est:79e01.s1 /CLONE=IMAGE:744216 /UG=Hs.116840 ESTs	NM_014498	1.70
gogi phosphoprotein 4	NM_020157	1.70
otorepelin	AF141392	1.70
RAN binding protein 7	AW238632	1.70
KIAA1098 protein	AK026548	1.70
KIAA0068 protein	AK025552	1.70
Consensus includes gb:AK025552.1 /DEF=Homo sapiens cDNA; FLJ21895 fls, clone HEP03467. /FEA=mRNA /DB_XREF=gi:10438101 /UG=Hs.306815 Homo sapiens cDNA; FLJ21895 fls, clone HEP03467	AK026120	1.70
KIAA0460 protein	AC005255	1.70
olfactory receptor, family 7, subfamily A, member 17	Y15014	1.70
UDP-Gal:beta-GlcNAc beta 1,3-galactosyltransferase, polypeptide 2	BF438282	1.70
PDZ domain containing guanine nucleotide exchange factor(GEF)1	BF058643	1.70
EGF-like repeats and discoidin-like domains 3	NM_000726	1.70
calcium channel, voltage-dependent, beta 4 subunit	M80927	1.70
chitinase 3-like 1 (cartilage glycoprotein-39)	M24900	1.70
nuclear receptor subfamily 1, group D, member 1	BC003541	1.70
hypothetical protein FLJ10488	AL049215	1.70
Dystonia musculorum of mouse, human homolog of	L25275	1.70
sulfotransferase family, cytosolic, 1A, phenol-preferring, member 3	NM_017488	1.70
adducin 2 (beta)	AB028816	1.70
phosphodiesterase 10A	R85779	1.69
spectrin, beta, non-erythrocytic 1	AL049394	1.69
hypothetical protein FLJ10648	NM_004108	1.69
ficollin (collagen/vibroninogen domain-containing lectin) 2 (hu collagen)	AW301305	1.69
KIAA1096 protein	AB088958	1.69
KIAA0870 protein	NM_031275	1.69
testis expressed sequence 12	AL022165	1.69
carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7	AL137592	1.69
dual oxidase 1	NM_025057	1.69
hypothetical protein FLJ23169	NM_023613	1.69
KIAA1009 protein	AW874823	1.69
Consensus includes gb:AW874823 /FEA=EST /DB_XREF=gi:8166026 /DB_XREF=est:EST386928 /UG=Hs.105411 ESTs	NM_003555	1.69
olfactory receptor, family 1, subfamily G, member 1	NM_004407	1.69
dentin matrix acidic phosphoprotein	BE563442	1.69
interleukin 1 receptor antagonist	NM_014289	1.69
calpain 6	AL134489	1.69
ribonuclease P (38kD)	X56687	1.69
upstream binding transcription factor, RNA polymerase I		

KIA0808 gene product
hypothetical protein FLJ20519
membrane-spanning 4-domains, subfamily A, member 4A
transcription factor binding to IGHM enhancer 3
v-rat-1 murine leukemia viral oncogene homolog 1
gb:L34164.1 /DEF=Human Ig rearranged mu-chain gene VH3-D2110-JH2, complete cds., polyreactive. /FEA=mRNA /DB_XREF=gi:576594 /FL=gb:L34164.1
muscle, skeletal, receptor tyrosine kinase
glutamate receptor, metabotropic 8
Consensus includes gb:AL137063 /DEF=Human DNA sequence from clone RP1-104L14 on chromosome 6 Contains part of the AKAP7 (A kinase (PRKA) anchor protein 7
KIA0451 gene product
chromobox homolog 8
endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform
inulin receptor substrate 1
lysosomal-associated membrane protein 2
SET translocation (myeloid leukemia-associated)
epithelial membrane protein 1
protocadherin 9
cyclin receptor-like factor 3
Escherichia coli /REF=J04423 /DEF=E coli bioD gene dehydrobiotin synthetase corresponding to nucleotides 5312-5559 of J04423 /LEN=676 (-5 and -3 represent transcript r
hypothetical protein
Consensus includes gb:BG433489 /FEA=EST /DB_XREF=gi:1333995 /DB_XREF=est:802497919F1 /CLONE=IMAGE:4611460 /UG=Hs.297257 ESTs
peptidyl arginine deiminase, type V
Consensus includes gb:AL121552 /DEF=Human DNA sequence from clone RP11-28009 on chromosome 20. Contains an RNF11 (ring finger protein 11) (SID1669) pseud
short stature homeobox 2
proteoglycan 1, secretory granule
NCK-associated protein 1
hypocretin (orexlin) receptor 1
interleukin 6 receptor
malignant fibrous histiocytoma amplified sequence 1
oligodendrocyte lineage transcription factor 2
VASA protein
aldehyde dehydrogenase 3 family, member B1
KIA0189 gene product
DKFZP586P0123 protein
core-binding factor, runt domain, alpha subunit 2; translocated to, 1; cyclin D-related
protein phosphatase 1, regulatory (inhibitor) subunit 6
hypothetical protein FLJ12457
suppression of tumorigenicity 14 (colon carcinoma, matrilysin, epithin)
Consensus includes gb:AK025094.1 /DEF=Homo sapiens cDNA: FLJ21441 fls, clone COL04422. /FEA=mRNA /DB_XREF=gi:10437541 /UG=Hs.269928 Homo sapiens cC AK025094
Consensus includes gb:AL049252.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564D193 (from clone DKFZp564D193). /FEA=mRNA /DB_XREF=gi:4499993 /UG=Hs.3020 AL049252
Consensus includes gb:AW190208 /FEA=EST /DB_XREF=gi:8464688 /DB_XREF=est:u60a12.x1 /CLONE=IMAGE:2679070 /UG=Hs.233789 ESTs
uncharacterized bone marrow protein BM029
forhead box D2
novel putative protein similar to YIL091C yeast hypothetical 84 kD protein from SG41-KTR7
Consensus includes gb:AK022408.1 /DEF=Homo sapiens cDNA FLJ12346 fls, clone MAMMA1002287, highly similar to Homo sapiens mRNA for Rab6 GTPase activating f
Consensus includes gb:AW813387 /FEA=EST /DB_XREF=gi:7318573 /DB_XREF=est:th71604.x1 /CLONE=IMAGE:2968254 /UG=Hs.278541 ESTs, Moderately similar to
suppressor of actin 1
Consensus includes gb:AK00145.1 /DEF=Homo sapiens cDNA FLJ20138 fls, clone COL07172. /FEA=mRNA /DB_XREF=gi:7020041 /UG=Hs.306384 Homo sapiens cDN AK00145
lectin, galactose-binding, soluble, 8 (galactin 8)
zinc finger protein 42 (myeloid-specific retinoic acid- responsive)
kruppel-related zinc finger protein hckrox
Consensus includes gb:AW150085 /FEA=EST /DB_XREF=gi:6197971 /DB_XREF=est:sg48a10.x1 /CLONE=IMAGE:2630779 /UG=Hs.271957 ESTs
hypothetical protein FLJ20084
PRO0650 protein
mucin 5, subtype B, tracheobronchial
arylsulfatase D
hypothetical protein FLJ23081
KIA0832 protein
Consensus includes gb:AK025077.1 /DEF=Homo sapiens cDNA: FLJ21424 fls, clone COL04157. /FEA=mRNA /DB_XREF=gi:10437519 /UG=Hs.287662 Homo sapiens cC AK025077
myelin protein zero-like 1
gb:U43279.1 /DEF=Human nucleoporin nup 38 mRNA, complete cds. /FEA=mRNA /PROD=nup 38 /DB_XREF=gi:1565299 /FL=gb:U43279.1
Consensus includes gb:AL050329 /DEF=Human DNA sequence from clone RP1-2217 on chromosome 6q11.1-12. Contains the 3 end of the gene KIA0244, the 3 end of AL050329

sialyltransferase 8 (alpha-2, 8-sialyltransferase) B
 melanoma cell adhesion molecule
 fibroblast growth factor 17
 solute carrier family 1 (glial high affinity glutamate transporter), member 2
 KIAA0683 gene product
 HSKM-B protein
 M11187 Human 28S rRNA sequence, length 5025 bases, 3 prime target bases 3331-4985
 hydridiform mole associated and imprinted
 zinc finger protein 193
 Consensus includes gb:AK022045.1 /DEF=Homo sapiens cDNA FLJ11983 fls, clone HEMBB1001337 /FEA=mRNA /DB_XREF=gi:10433364 /UG=Hs.193162 Homo sapiens cDNA AW173157 /FEA=EST /DB_XREF=gi:8439105 /DB_XREF=est:84612.x1 /CLONE=IMAGE:2663502 /UG=Hs.26860 Homo sapiens mRNA; cDNA AW173157
 glycerol-3-phosphate dehydrogenase 1 (soluble)
 hemoglobin, gamma G
 villiglo-associated protein VIT-1
 protein kinase, interferon-inducible double stranded RNA dependent
 zinc finger protein 223
 hypothetical protein PRO2714
 aldehyde oxidase 1
 hepatocyte nuclear factor 4, alpha
 MAD, mothers against decapentaplegic homolog 5 (Drosophila)
 hypothetical protein MGC2508
 dynein, cytoplasmic, intermediate polypeptide 2
 insulin-like growth factor binding protein 2 (38kD)
 hypocretin (orexin) receptor 2
 hypothetical protein FLJ22792
 disc, large (Drosophila) homolog 1
 beta-site APP-cleaving enzyme
 ATPase, H⁺ transporting, lysosomal (vacuolar proton pump) 9kD
 v-akt murine thymoma viral oncogene homolog 1
 transcription factor (p38 interacting protein)
 topoisomerase (DNA) I
 hypothetical protein FLJ13340
 RNA binding motif protein 12
 placental growth factor, vascular endothelial growth factor-related protein
 EST-YD1 protein
 putative X-linked retinopathy protein
 lethal (3) malignant brain tumor (3)mbt protein (Drosophila) homolog
 aquaporin 1 (channel-forming integral protein, 28kD)
 KIAA0841 protein
 hypothetical gene CG018
 lysosomal-associated membrane protein 3
 solute carrier family 1 (glutamate transporter), member 7
 KIAA0908 protein
 aldehyde dehydrogenase 3 family, member B2
 protease, serine, 12 (neutrypsin, molopsin)
 KIAA0528 gene product
 KIAA0875 gene product
 manic fringe homolog (Drosophila)
 retinoic acid receptor, beta
 basement membrane-induced gene
 von Hippel-Lindau syndrome
 general transcription factor IIIC, polypeptide 2 (beta subunit, 110kD)
 insulin-like growth factor binding protein 5
 interleukin 1, zeta
 cytochrome P450, subfamily IIA (nifedipine oxidase), polypeptide 4
 Consensus includes gb:AB007970.1 /DEF=Homo sapiens mRNA, chromosome 1 specific transcript KIAA0501, /FEA=mRNA /DB_XREF=gi:3413945 /UG=Hs.223020 Hom
 chymotrypsinogen B1
 Sp3 transcription factor
 gamma-glutamyl carboxylase
 Consensus includes gb:AK024553.1 /DEF=Homo sapiens cDNA: FLJ20900 fls, clone ADKA03593, /FEA=mRNA /DB_XREF=gi:10436861 /UG=Hs.30687 Homo sapiens c
 Consensus includes gb:AK025072.1 /DEF=Homo sapiens cDNA: FLJ21419 fls, clone COL04084, /FEA=mRNA /DB_XREF=gi:10437512 /UG=Hs.306759 Homo sapiens cD
 zinc finger protein 238

NM_006011 1.68
 M29277 1.68
 NM_003887 1.68
 NM_004171 1.68
 AL137394 1.67
 AB09870 1.67
 M11187 1.67
 AF241534 1.67
 NM_006299 1.67
 AK022045 1.67
 cDNA AW173157 1.67
 AI368018 1.67
 NM_000184 1.67
 NM_018693 1.67
 NM_002759 1.67
 NM_013361 1.67
 NM_018534 1.67
 AB046692 1.67
 Z49825 1.67
 AF010801 1.67
 AA639585 1.67
 AF250307 1.67
 NM_000597 1.67
 NM_001526 1.67
 NM_024821 1.67
 BC251175 1.67
 NM_012104 1.67
 BE043477 1.67
 NM_005163 1.67
 NM_017569 1.67
 AK000861 1.67
 AW025108 1.67
 NM_025085 1.67
 AL514547 1.67
 BC001422 1.67
 NM_021208 1.67
 S58722 1.67
 AW445040 1.67
 AL518391 1.67
 AA845355 1.67
 AL049786 1.67
 NM_014398 1.67
 NM_006671 1.67
 AB87102 1.66
 NM_000685 1.66
 AB10787 1.66
 AB011100 1.66
 NM_014648 1.66
 AF760053 1.66
 X04014 1.66
 AB035482 1.66
 NM_000551 1.66
 NM_001521 1.66
 M65062 1.66
 NM_014439 1.66
 NM_017460 1.66
 AB007870 1.66
 NM_001906 1.66
 AU145005 1.66
 BE326952 1.66
 AK024553 1.66
 Consensus includes gb:AK025072 1.66
 AK000847 1.66

hypothetical protein FLJ13605
 Consensus includes gb:AU145680 /FEA=EST /DB_XREF=gi:11007201 /DB_XREF=est:AU145680 /CLONE=HEMBA1005500 /UG=Hs.306613 Homo sapiens cDNA FLJ111372
 hypothetical protein MGC5528
 hypothetical protein KIAA1102 protein
 lyxuretinase (L-xyuretinase hydrolase)
 muscle-specific beta 1 integrin binding protein
 chromosome 21 open reading frame 2
 hypothetical protein PRO1584
 neuromedin B receptor
 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit e
 actin-like protein-like 1 (Drosophila)
 protein phosphatase 2, regulatory subunit B (856), alpha isoform
 transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)
 KIAA0680 gene product
 KIAA0414 protein
 Consensus includes gb:AW299245 /FEA=EST /DB_XREF=gi:6708922 /DB_XREF=est:xs48d07.x1 /CLONE=IMAGE:2772877 /UG=Hs.62601 Homo sapiens mRNA; cDNA
 metapase chromosome protein 1
 protocadherin alpha 9
 hypothetical protein FLJ10044
 potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3
 transcription factor 7 (T-cell specific, HMG-box)
 mitogen-activated protein kinase 8 interacting protein 2
 chemokine (C-C motif) receptor 2
 Consensus includes gb:BF434424 /FEA=EST /DB_XREF=gi:11446712 /DB_XREF=est:768608.x1 /CLONE=IMAGE:3844393 /UG=Hs.107184 spectrin, beta, non-erythrocyte
 Consensus includes gb:AU147942 /FEA=EST /DB_XREF=gi:11009463 /DB_XREF=est:AU147942 /CLONE=MAMMA1002188 /UG=Hs.287517 Homo sapiens cDNA FLJ1212
 olfactory receptor, family 10, subfamily H, member 1
 Consensus includes gb:BC004145.1 /DEF=Homo sapiens, clone IMAGE:3509783, mRNA, partial cds, /FEA=mRNA /PROD=Unknown (protein for IMAGE:3509783) /DB_XI BC004145
 serine/threonine kinase 11 (Peutz-Jeghers syndrome)
 RAB31, member RAS oncogene family
 chromosome 2 open reading frame 3
 thyrotropin-releasing hormone degrading ectoenzyme
 pax6, alpha
 tumor endothelial marker 7 precursor
 flavin containing monooxygenase 2
 gastrulation brain homeo box 2
 thrombospondin 1
 Cbort10 protein
 hypothetical protein FLJ12121
 tumor necrosis factor (ligand) superfamily, member 13
 KIAA0339 gene product
 Cluster Ind. AL050186 Homo sapiens mRNA; cDNA DKFZp586D1122 (from clone DKFZp586D1122) /cds=UNKNOWN /gb=AL050166 /ig=4884381 /ug=Hs.26295 /len=26; N30342
 MAX binding protein
 integrin, alpha 2b (platelet) glycoprotein IIb of fibrin complex, antigen CD41B)
 SREBP CLEAVAGE-ACTIVATING PROTEIN
 inhibitor, beta B (ecton AB beta polypeptide)
 O-substrate
 LIM domain protein
 enhancer of zeste homolog 1 (Drosophila)
 glutamate receptor, metabotropic 1
 AND-1 protein
 microtubule-associated protein 1A
 arachidonate 12-lipoxygenase
 purine-rich element binding protein G
 lamin A/C
 thyroid hormone receptor interactor 6
 KIAA0870 protein
 myelin-associated oligodendrocyte basic protein
 Consensus includes gb:AL117643.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M245 (from clone DKFZp434M245) /FEA=mRNA /DB_XREF=gi:5912233 /UG=Hs.5286 /AL117643
 BAI1-associated protein 3
 thyroid transcription factor 1
 hypothetical protein PRO2007
 immunoglobulin heavy constant gamma 3 (G3m marker)

transferrin
Consensus includes gb:AU147962 /FEA=EST /DB_XREF=gi:11009483 /DB_XREF=est:AU147962 /CLONE=MAMMA1002250 /UG=Hs.301509 Homo sapiens cDNA FLJ12
1.64

RelA-associated inhibitor
NM_006663
1.64

dimethylarginine dimethylaminohydrolase 1
AL078459
1.64

metal-regulatory transcription factor 1
AW182367
1.64

heterogeneous nuclear ribonucleoprotein H3 (2H9)
L32610
1.64

cyokeratin 2
NM_015848
1.64

solute carrier family 28 (sulfate transporter), member 2
AI025519
1.64

adaptor-related protein complex 1, sigma 1 subunit
BC003561
1.64

phospholipase A2 receptor 1, 180kD
U17033
1.64

ACO for serine protease homologue
NM_017509
1.64

hypothetical protein FLJ12868
NM_024997
1.64

Consensus includes gb:W28849 /FEA=EST /DB_XREF=gi:1308815 /DB_XREF=est:52g6 /UG=Hs.33862 ESTs
W28849
1.64

PRO0097 protein
NM_014114
1.64

HCNP protein; XPA-binding protein 2
NM_020196
1.64

neurophilin 3
AI837333
1.64

solute carrier family 6 (neurotransmitter transporter, laurine), member 6
BC006252
1.64

BRF2, subunit of RNA polymerase III transcription initiation factor. BRF1-like
AF288153
1.64

Consensus includes gb:AL121903 /DEF=Human DNA sequence from clone RP1-155G6 on chromosome 20 Contains part of the gene for briefeldin A-inhibited guanine nud
AL121903
1.64

BH-protocadherin (brain-heart)
AB006757
1.64

HYA22 protein
BF002474
1.64

zinc finger protein 175
NM_007147
1.64

F-box only protein 9
AL137520
1.64

CAMP response element-binding protein CRE-BPa
NM_004904
1.64

NBR2
NM_005821
1.64

ATP-binding cassette, sub-family A (ABC1), member 1
NM_005502
1.64

SON DNA binding protein
AI836458
1.64

Phosphatidyglycerophosphate Synthase
NM_024419
1.64

CGI-43 protein
NM_015622
1.64

Escherichia coli /REF=J04423 /DEF=E coli bioD gene dehydrobiotin synthetase corresponding to nucleotides 5024-5244 of J04423 /LEN=876 (-5 and -3 represent transcript r
J04423
1.64

tachykinin receptor 1
M81797
1.64

activating transcription factor 1
AA34345
1.64

Consensus includes gb:AL078638.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 117929. /FEA=mRNA /DB_XREF=gi:5051866 /UG=Hs.227171
AL078638
1.63

S100 calcium binding protein, beta (neural)
BC001766
1.63

pancreatic polypeptide 2
NM_021092
1.63

Consensus includes gb:AU147200 /FEA=EST /DB_XREF=gi:11008721 /DB_XREF=est:AU147200 /CLONE=HEMBB1002699 /UG=Hs.296698 Homo sapiens cDNA FLJ12
AU147200
1.63

homonally upregulated Neu-associated kinase
NM_014586
1.63

guanine nucleotide binding protein (G protein), beta 5
NM_016194
1.63

F-box only protein 22
NM_012170
1.63

Consensus includes gb:AF070602.1 /DEF=Homo sapiens clone 24504 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387982 /UG=Hs.51649 Homo sapiens clone 24504 /AF070602
AF070602
1.63

Consensus includes gb:AK022886.1 /DEF=Homo sapiens cDNA FLJ12624 fls. clone NT2RM4001754. /FEA=mRNA /DB_XREF=gi:10434225 /UG=Hs.306639 Homo sapien
AK022886
1.63

HIV-1 Inducer of short transcripts binding protein: lymphoma related factor
AW027070
1.63

mitochondrial solute carrier
NM_018579
1.63

collagen, type VIII, alpha 1
BE877796
1.63

likely homolog of rat and mouse retinoid-inducible serine carboxypeptidase
NM_021626
1.63

nuclear receptor coactivator 1
AI872428
1.63

VW domain binding protein 4 (formin binding protein 21)
NM_007187
1.63

gb:NM_018634.1 /DEF=Homo sapiens hypothetical protein PRO2893 (PRO2893), mRNA. /FEA=mRNA /GEN=PRO2893 /PROD=hypothetical protein PRO2893 /DB_XREF
NM_018634
1.63

ribosomal protein S8 kinase, 70kD, polypeptide 2
NM_003952
1.63

muscle, skeletal, receptor tyrosine kinase
NM_005592
1.63

purine-rich element binding protein A
BE222739
1.63

Kruppel-like factor 4 (gut)
NM_004235
1.63

Wiskott-Aldrich syndrome (eczema-thrombocytopenia)
NM_000377
1.63

aldehyde dehydrogenase 1 family, member A2
NM_003888
1.63

Consensus includes gb:AL108696.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 21920. /FEA=mRNA /DB_XREF=gi:5689771 /UG=Hs.185701 /AL108696
AL108696
1.63

Consensus includes gb:AW451230 /FEA=EST /DB_XREF=gi:8992006 /DB_XREF=est:UH-H-B13-ah-a-01-Q-U1.s1 /CLONE=IMAGE:2736649 /UG=Hs.257332 ESTs. Highly
AW451230
1.63

fibronectin leucine rich transmembrane protein 3
NM_0213281
1.63

phosphotyrosine phosphatidylserine inorganic pyrophosphate phosphatase
NM_022126
1.63

chondroitin sulfate proteoglycan 4 (melanoma-associated)
NM_001897
1.63

arrestin A2 pseudogene 1
M82896
1.63

Consensus includes gb:AL121981 /DEF=Human DNA sequence from clone RP5-1061C18 on chromosome 1p36.22-36.33. Contains ESTs, STSs, GSSs and a CoQ island.
AL121981
1.63

sterile-alpha motif and leucine zipper containing kinase AZK
NM_016653
1.63

DKF ZP586J0119 protein
AF112207
1.63

hypothetical protein FLJ11898	NM_024976	1.62
hepatitis delta antigen-interacting protein A	BE9868372	1.63
hypothetical protein dJ79617.1	NM_052225	1.63
slalophorin (gpl.115, leukosialin, CD43)	NM_003123	1.63
ras-like protein	AW771590	1.63
KIA0523 protein	AB011095	1.63
hypothetical protein FLJ13590	NM_024840	1.63
hypothetical protein dJ465N24.2.1	AF287858	1.62
KIAA1100 protein	NM_014801	1.62
cadherin 19, type 2	NM_021153	1.62
lunatic fringe homolog (Drosophila)	U94354	1.62
methyl-CpG binding domain protein 2	A1827820	1.62
programmed cell death 2	A4784868	1.62
canilaine peptidyltransferase 1, liver	BF001714	1.62
KIAA0376 protein	AL117532	1.62
Consensus includes gb:AL110201.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586F1622 (from clone DKFZp586F1622) /FEA=mrna /DB_XREF=gi:5817120 /UG=Hs.271	AL110201	1.62
stathmin-like 4	NM_030795	1.62
hypothetical protein FLJ21269	NM_025107	1.62
synaptotagmin 1	AV723167	1.62
adrenergic, beta-3-, receptor	X70812	1.62
DnaJ (Hsp40) homolog, subfamily B, member 1	NM_006145	1.62
neural retina leucine zipper	M81840	1.62
hair/enhancer-of-split related with YRPW motif 1	R81374	1.62
Consensus includes gb:U92027.1 /DEF=Human clone 61501 defective mariner transposon Hsma2 mRNA sequence. /FEA=mrna /DB_XREF=gi:2052469 /UG=Hs.15921	U92027	1.62
amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	NM_000484	1.62
homoio box D13	NM_000523	1.62
transcription factor 20 (AR1)	AB006630	1.62
hypothetical protein FLJ14326	AK024388	1.62
hypothetical protein FLJ13710	NM_024817	1.62
zinc finger protein 272	X78931	1.62
related to the N terminus of tre	AF034374	1.62
Consensus includes gb:AK025580.1 /DEF=Homo sapiens polydendrum cofactor biosynthesis protein C mRNA, complete cds. /FEA=mrna /PI	NM_014688	1.62
potassium channel, subfamily K, member 7	AK025580	1.62
Consensus includes gb:AB55015 /FEA=EST /DB_XREF=gi:4738994 /DB_XREF=est:wb66a10.x1 /CLONE=IMAGE.2310618 /UG=Hs.3843 Homo sapiens mRNA, cDNA DI	NM_005714	1.62
UV-B repressed sequence, HUR 7	BE148534	1.62
Consensus includes gb:AL050026.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586C0224 (from clone DKFZp586C0224) /FEA=mrna /DB_XREF=gi:4884285 /UG=Hs.30	AL050026	1.62
glutamate decarboxylase 1 (brain, 67KD)	NM_013445	1.62
chromosome 1 open reading frame 6	AK023354	1.62
c-myc binding protein	D50892	1.62
transcription factor 20 (AR1)	U19345	1.62
karyopherin alpha 1 (importin alpha 5)	BC002374	1.62
NIMA (never in mitosis gene a)-related kinase 2	NM_002497	1.62
hypothetical protein FLJ23168	NM_024628	1.62
collagen, type VI, alpha 1	M20776	1.62
desmin	AK022087	1.62
hypothetical protein MGC5178	AI580112	1.62
casein kinase 1, gamma 2	NM_001319	1.62
Slah-interacting protein	BC005975	1.62
hypothetical protein FLJ21369	NM_024802	1.62
hypothetical protein FLJ11588	NM_024603	1.62
Rhesus blood group-associated glycoprotein	AF178841	1.62
KIAA1538 protein	BC003177	1.62
KIAA0557 protein	BC434168	1.62
Rad50-interacting protein 1	AB011129	1.62
Consensus includes gb:S65921.1 /DEF=anti-colorectal carcinoma light chain=glycoprotein CANAG-50 specific IgG1 kappa human, 19.9 hybridoma, antibody 1118NS19.9.1	NM_021930	1.62
keratin 5 (epidermolysis bullosa simplex, Dowling-Mearns/Kobner/Weber-Cockayne types)	NM_000424	1.62
Consensus includes gb:AK024602.1 /DEF=Homo sapiens cDNA: FLJ20949 /fis, clone ADSE01902 /FEA=mrna /DB_XREF=gi:10436915 /UG=Hs.220255 Homo sapiens c	AK024602	1.62
serine/threonine kinase 6	NM_003158	1.61
interleukin 13 receptor, alpha 1	U62858	1.61
ets variant gene 6 (TEL oncogene)	AF041811	1.61
limodulin 1 (phosphorylase kinase, delta)	M27319	1.61

KIAA0138 gene product 1.61
aldehyde dehydrogenase 1 family, member A2 NM_014649 1.61
heat shock 70kD protein 2 AB013226 1.61
hypothetical gene FLJ00060 U58725 1.61
7-dehydrocholesterol reductase AK024467 1.61
leucine zipper, down-regulated in cancer 1 NM_001360 1.61
karyopherin beta 2b, transportin NM_012317 1.61
WW domain-containing oxidoreductase AK022217 1.61
cell division cycle 25B NM_016373 1.61
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson relaxin 2 (H2) AL109804 1.61
synaptotagmin XIII NM_022969 1.61
hypothetical protein MY014 NM_005059 1.61
like mouse brain protein E46 AW089736 1.61
putative nuclear protein NM_030818 1.61
mitogen-activated protein kinase kinase 3 Z95331 1.61
ubiquitin-conjugating enzyme E2B (RAD6 homolog) AA046650 1.61
Consensus includes gb:AF052160.1 /DEF=Homo sapiens clone 24629 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360471 /UG=Hs.142570 Homo sapiens clone 24629 A768723 1.61
angiotensin-like 2 NM_003618 1.61
SEC22 vesicle trafficking protein-like 1 (S. cerevisiae) NM_023938 1.61
tumor necrosis factor (ligand) superfamily, member 13 D13413 1.61
ATPase, Ca++ transporting, plasma membrane 2 NM_004102 1.61
hypothetical protein MGC2742 NM_004102 1.61
heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A) NM_004102 1.61
fatty acid binding protein 3, muscle and heart (mammary-derived growth inhibitor) NM_004102 1.61
Consensus includes gb:AF319523.1 /DEF=Homo sapiens RT-L1 mRNA, complete sequence. /FEA=mRNA /PROD=RT-L1 /DB_XREF=gi:13310144 /UG=Hs.326775 Homo sapiens clone 24629 AF319523 1.61
UDP-Gal-4-epiChNAc beta 1,3-galactosyltransferase, polypeptide 5 NM_006057 1.61
J04423 E coli bioD gene deshydrobiotin synthetase (-5 and -3 represent transcript regions 5 prime and 3 prime respectively) AF114013 1.61
ATPase, Ca++ transporting, plasma membrane 2 J04423 1.61
Interleukin 1 receptor accessory protein R52647 1.61
Consensus includes gb:AW293343 /FEA=EST /DB_XREF=gi:6699905 /DB_XREF=est:U1-H-B12-ah1-d-05-o-U1.s1 /CLONE=IMAGE:2726841 /UG=Hs.283978 Homo sapiens clone 24629 AF167070 1.61
LIM domain kinase 1 NM_023314 1.61
ATP-binding cassette, sub-family B (MOT7AP), member 11 NM_002314 1.61
dipeptidylpeptidase IV (CD26, adenosine deaminase complexing protein 2) NM_003742 1.61
musculin (activated B-cell factor-1) M80536 1.61
hypothetical protein MGC3489 AF060154 1.61
KIAA0042 gene product NM_024055 1.61
KIAA1334 protein NM_014875 1.61
CGI-148 protein AA460694 1.61
ubiquitin protein ligase E3A (human papilloma virus E6-associated protein, Angelman syndrome) NM_016078 1.61
Consensus includes gb:AL096727.1 /DEF=Homo sapiens mRNA: cDNA DKFZP434B104 (from clone DKFZP434B104). /FEA=mRNA /DB_XREF=gi:5419860 /UG=Hs.1682 U84404 1.61
aldehyde dehydrogenase 7 family, member A1 BC002515 1.61
APMCF1 protein NM_021203 1.61
purinergic receptor P2X, ligand-gated ion channel, 4 NM_002560 1.61
Consensus includes gb:AK023174.1 /DEF=Homo sapiens cDNA: FLJ11521 fs, clone COL05880. /FEA=mRNA /DB_XREF=gi:10437836 /UG=Hs.306777 Homo sapiens clone 24629 AL163248 1.61
zinc finger protein 284 NM_012269 1.61
hyaluronoglucosaminidase 4 NM_005181 1.61
carbonic anhydrase III, muscle specific W87468 1.61
hypothetical protein FLJ20783 BC002511 1.61
carbonyl reductase 1 A344075 1.61
gamma-glutamyltransferase 2 NM_001148 1.60
ankyrin 2, neuronal NM_023942 1.60
hypothetical protein MGC3038 U73191 1.60
potassium inwardly-rectifying channel, subfamily J, member 15 NM_001330 1.60
cardiotrophin 1 M28095 1.60
calcitonin/calcitonin-related polypeptide, alpha BF197222 1.60
Consensus includes gb:BF197222 /FEA=EST /DB_XREF=gi:11085906 /DB_XREF=est:7m8b07.x1 /CLONE=IMAGE:3561949 /UG=Hs.12342 Homo sapiens clone 24538 BF197222 1.60
CD5 antigen-like (scavenger receptor cysteine rich family) NM_005894 1.60
CGI-109 protein AF151867 1.60
cyclochrome P450, subfamily VIIIB (cytochrome P450) NM_004820 1.60
Consensus includes gb:AK001448.1 /DEF=Homo sapiens cDNA: FLJ10586 fs, clone NT2RP2003986. /FEA=mRNA /DB_XREF=gi:7022713 /UG=Hs.301255 Homo sapiens clone 24538 AK001448 1.60
glutamate receptor, metabotropic 5 NM_000842 1.60
preprorelactin-releasing peptide NM_015893 1.60

ectopic viral integration site 5
 N-acylsphingosine amidohydrolase (acid ceramidase)
 Consensus includes gb:BG230612 /FEA=EST /DB_XREF=gi:12725854 /DB_XREF=astna141b09.x1 /CLONE=IMAGE:4143329 /UG=Hs.13998 Homo sapiens cDNA: FLJ2: BG230612
 RBP1-like protein
 signal transducing adaptor molecule (SH3 domain and ITAM motif) 2
 hypothetical protein FLJ20856
 CED-6 protein
 zinc finger protein 76 (expressed in testis)
 cellular retinoic acid binding protein 2
 Consensus includes gb:U78287.1 /DEF=Human clone 23589 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710280 /UG=Hs.11506 Human clone 23589 mRNA sequence U79297
 Consensus includes gb:AA700015 /FEA=EST /DB_XREF=gi:2702978 /DB_XREF=est169c11.s1 /CLONE=IMAGE:436052 /UG=Hs.287995 Homo sapiens cDNA: FLJ2318 AA700015
 GTP binding protein overexpressed in skeletal muscle
 KIAA0416 protein
 thrombospondin 2
 growth arrest-specific 2
 Consensus includes gb:AA653456 /FEA=EST /DB_XREF=gi:2589627 /DB_XREF=estag66c08.s1 /CLONE=IMAGE:1127918 /UG=Hs.270495 ESTs, Weakly similar to ubiq
 solute carrier family 2 (facilitated glucose transporter), member 3
 Cyclin E1
 KIAA1451 protein
 Microfibril-associated glycoprotein-2
 sodium channel, voltage-gated, type IV, alpha polypeptide
 Consensus includes gb:R78289 /FEA=EST /DB_XREF=gi:853409 /DB_XREF=est1y81n12.s1 /CLONE=IMAGE:145703 /UG=Hs.287419 Homo sapiens cDNA: FLJ11505 fts R78289
 hypothetical protein FLJ12681
 apolipoprotein A-II
 perlecan
 KIAA0987 protein
 hypothetical protein PRO2435
 Consensus includes gb:AA491625 /FEA=EST /DB_XREF=gi:2221187 /DB_XREF=estne80d01.s1 /CLONE=IMAGE:10561 /UG=Hs.293940 ESTs
 S100 calcium binding protein A12 (calgranulin C)
 keratin 17
 hemochromatosis
 annexin A5
 synaptotagmin 1
 chromosome 8 open reading frame 10
 basic leucine zipper nuclear factor 1 (JEM-1)
 activating transcription factor 5
 gb:BC006361.1 /DEF=Homo sapiens, clone MGC:13137, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:13137) /DB_XREF=gi:13823508 /FL=gb:BC006361
 gb:U35622.2 /DEF=Homo sapiens EWS proteinE1A enhancer binding protein chimera mRNA, complete cds. /FEA=mRNA /PROD=EWS proteinE1A enhancer binding prot U35622
 retinoblastoma binding protein 1
 solute carrier family 4, sodium bicarbonate cotransporter, member 4
 estrogen receptor 1
 hypothetical protein FLJ10461
 neighbor of A-kinase anchoring protein 95
 FYN binding protein (FYN-120130)
 hypothetical protein MGC5306
 skin-specific protein
 gb:AF202635.1 /DEF=Homo sapiens PP1200 mRNA, complete cds. /FEA=mRNA /PROD=PP1200 /DB_XREF=gi:10732645 /UG=Hs.302135 Homo sapiens PP1200 mRNA AF202635
 hypothetical protein PRO1438
 G protein-coupled receptor
 A kinase (PKA) anchor protein 6
 GDNF family receptor alpha 1
 amyloid beta (A4) precursor-like protein 2
 fibroblast growth factor 12
 hypothetical protein DKFZp761G1913
 KIAA0346 protein
 vav 2 oncogene
 hypothetical protein FLJ20004
 2.19 gene
 Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 2
 aggrecan 1 (chondroitin sulfate proteoglycan, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122)
 formyl peptide receptor 1
 KIAA1025 protein
 KIAA0669 gene product

1.60 NM_005665
 1.60 U4764
 1.60 BG230612
 1.60 AL133600
 1.60 NM_025143
 1.60 AK023668
 1.60 NM_003427
 1.60 NM_001878
 1.60 U79297
 1.60 AA700015
 1.60 NM_005261
 1.60 NM_015564
 1.60 NM_003247
 1.60 NM_005256
 1.60 AA653456
 1.60 NM_006931
 1.60 AI671049
 1.60 BF970829
 1.60 AW665892
 1.60 NM_000334
 1.60 R78289
 1.60 NM_005621
 1.60 Z19574
 1.60 AF150864
 1.60 NM_001154
 1.60 NM_003895
 1.60 NM_008781
 1.60 NM_003666
 1.60 AF101388
 1.60 BC006361
 1.60 U35622
 1.60 NM_002892
 1.60 AF011390
 1.59 AI073549
 1.59 NM_018098
 1.59 AB016332
 1.59 AF198052
 1.59 NM_024116
 1.59 NM_014357
 1.59 AF202635
 1.59 NM_018600
 1.59 U73531
 1.59 NM_004274
 1.59 NM_005264
 1.59 BC000373
 1.59 AL119322
 1.59 AL136557
 1.59 AA521267
 1.59 NM_003371
 1.59 NM_017616
 1.59 BC002934
 1.59 AF109161
 1.59 NM_013227
 1.59 NM_002029
 1.59 BG426889
 1.59 AF201291

colony stimulating factor 1 (macrophage)	M76453	1.58
Consensus includes gb:AL109667.1 /DEF=Homo sapiens mRNA (full length insert cDNA clone EUROMIMAGE 381867 /FEA=mRNA /DB_XREF=gi:5699815 /UG=Hs.124154	AL109667	1.58
N-myristoyltransferase 1	A1570834	1.58
Consensus includes gb:AL359578.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547N163 /FEA=mRNA /DB_XREF=gi:5855637 /UG=Hs.3065	AL359578	1.58
KIAA1693 protein	A869898	1.58
arachidonate 12-lipoxygenase, 12R type	NM_001139	1.58
contactin 2 (axonal)	NM_005076	1.58
neurofilin 2	AF280546	1.58
CD1E antigen, e polypeptide	NM_030893	1.58
butyrophilin, subfamily 3, member A3	NM_006994	1.58
HSPC135 protein	NM_014170	1.58
Consensus includes gb:AL049435.1 /DEF=Homo sapiens mRNA; cDNA DKFZp5880220 (from clone DKFZp5880220) /FEA=mRNA /DB_XREF=gi:4500218 /UG=Hs.171	AL049435	1.58
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12E47)	A655986	1.58
KIAA0480 gene product	AW299294	1.58
Consensus includes gb:A952772 /FEA=EST /DB_XREF=gi:5745082 /DB_XREF=est:wr50d05.x1 /CLONE=IMAGE:2491113 /UG=Hs.300865 Homo sapiens clone BCSynL	A952772	1.58
phosphodiesterase 1B, calmodulin-dependent	NM_000924	1.58
thymine-DNA glycosylase	BF674842	1.58
growth factor independent 1	NM_005263	1.58
interferon induced transmembrane protein 1 (9-27)	A4749101	1.58
carboxylesterase-related protein	NM_016280	1.58
guanine nucleotide binding protein (G protein), alpha activating activity polypeptide O	AA758795	1.58
rhoGac guanine nucleotide exchange factor (GEF) 2	NM_004723	1.58
KIAA0266 gene product	NM_014699	1.58
Consensus includes gb:U82306.1 /DEF=Homo sapiens unknown protein mRNA, partial cds /FEA=mRNA /PROD=unknown /DB_XREF=gi:1938333 /UG=Hs.123081 Homo	U82306	1.58
hypothetical protein FLJ21777	AK025430	1.58
DNA (cytosine-5-methyltransferase 3 beta	NM_006892	1.58
KIAA0475 gene product	NM_014864	1.58
vacuolar protein sorting 45A (yeast)	NM_007258	1.58
uncharacterized hematopoietic stem/progenitor cells protein MDS029	NM_018464	1.58
hypothetical protein FLJ21603	NM_024762	1.58
Pyruvate dehydrogenase complex, lipoyl-containing component X; E3-binding protein	NM_003477	1.58
guanylate kinase 1	AW182392	1.58
general transcription factor IIC, polypeptide 2 (beta subunit, 110kD)	AW194657	1.58
Consensus includes gb:AF257059 /DEF=Homo sapiens prothymosin alpha (PTMA) gene, complete cds /FEA=CDS /DB_XREF=gi:3037944 /UG=Hs.283947 Homo sapiens	AF257059	1.58
Consensus includes gb:AW134608 /FEA=EST /DB_XREF=gi:6138154 /DB_XREF=est:U1-H-B11-abo-e-04-0-UI.s1 /CLONE=IMAGE:2712654 /UG=Hs.168941 ESTs	AW134608	1.58
complement component (3B/4b) receptor 1, including Knops blood group system	X14362	1.58
cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4)	NM_001262	1.58
chromosome 16 open reading frame 44	NM_024731	1.58
Consensus includes gb:AA017093 /FEA=EST /DB_XREF=gi:1479257 /DB_XREF=est:ze37h10.s1 /CLONE=IMAGE:361219 /UG=Hs.271780 ESTs	AA017093	1.58
centaurin, alpha 2	NM_018404	1.58
TAF4A RNA polymerase II, TATA box binding protein (TBP)-associated factor, 135 kD	AI744029	1.58
secreted protein of unknown function	BF440021	1.58
tenilin AC	NM_005572	1.58
fizzled homolog 1 (Drosophila)	AF072872	1.58
discs, large (Drosophila) homolog-associated protein 1	NM_004746	1.58
neuronal specific transcription factor DAT1	AL050152	1.58
U2 small nuclear ribonucleoprotein auxiliary factor (65kD)	AB101556	1.58
Consensus includes gb:AK025065.1 /DEF=Homo sapiens cDNA: FLJ21412 fls. clone COL03987 /FEA=mRNA /DB_XREF=gi:10437504 /UG=Hs.288066 Homo sapiens cd	AK025065	1.58
myosin IC	BE790157	1.58
VLCS-H1 protein	NM_014031	1.58
flavin containing monooxygenase 2	AL021026	1.58
Interleukin 7 receptor	NM_002185	1.58
Nijmegen breakage syndrome 1 (nibrin)	NM_002485	1.58
aquaporin 6, kidney specific	AL137716	1.57
SRP (sex determining region Y)-box 9 (campomelic dysplasia, autosomal sex-reversal)	NM_000346	1.57
hypothetical protein FLJ21240	NM_024847	1.57
succinate-CoA ligase, ADP-forming, beta subunit	NM_003850	1.57
guanine nucleotide binding protein (G protein), alpha 15 (Gq class)	NM_002068	1.57
high-mobility group protein 2-like 1	AL078310	1.57
hypothetical protein FLJ10848	NM_018167	1.57
carbonic anhydrase VA, mitochondrial	NM_001739	1.57
translocated promoter region (to activated MET oncogene)	BF110993	1.57
hypothetical protein FLJ20473	NM_017838	1.57

	hypothetical protein LOC37158	NM_0290433
	Parkinson disease (autosomal recessive, juvenile) 2, parkin	NM_0045622
	hypothetical protein FLJ23168	NM_0250555
	nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100)	U08609
	micosomal triglyceride transfer protein (large polypeptide, 8kD)	A8RZ3321
	Consensus includes gb:AUI58608 /FEA=EST /DB_XREF=ggi:11020127 /DB_XREF=est:AU158606 /CLONE=PLACE3000406 /UG=Hs.287638 Homo sapiens cDNA FLJ143 AU158606	AF091627
	tumor protein 63 kDa with strong homology to p53	U08609
	Consensus includes gb:AP035317.1 /DEF=Homo sapiens clone 23892 mRNA sequence, /FEA=mRNA /DB_XREF=ggi:2861080 /UG=Hs.91916 Homo sapiens clone 23892, AF035317	AF035317
	PR domain containing 9	NM_020227
	caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)	U13699
	hypothetical protein FLJ10034	NM_017973
	piron protein (p27-30) [Creutzfeldt-Jakob disease, Gerstmann-Straussler-Scheinker syndrome, fatal familial Insomnia]	NM_000311
	karyopherin alpha 1 (importin alpha 5)	NM_002264
	neurotrophic tyrosine kinase, receptor, type 3	NM_002530
	T-cell acute lymphocytic leukemia 1	X51990
	transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	AW062341
	hypothetical protein FLJ10891	NM_018260
	actinin, alpha 1	AU126260
	Consensus includes gb:AL54245 /FEA=EST /DB_XREF=ggi:12894837 /DB_XREF=est:AL54245 /CLONE=CSD01082YA02 (S prime) /UG=Hs.288989 nuclear receptor sut AL54245	M95178
	myosin X	NM_012334
	hypothetical protein FLJ10460	NM_018097
	TAKF RNA polymerase II, TATA box binding protein (TBP)-associated factor, 150 kD	AK001618
	dihydropicolinate branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease)	NM_001918
	NTF2-related export protein 1	NM_013248
	hairless homolog (mouse)	AF039198
	solute carrier family 4, sodium bicarbonate cotransporter, member 7	AF047033
	Rat guanine nucleotide exchange factor RaiGPS1A	AF221098
	kallikrein 3, (prostate specific antigen)	NM_001648
	T cell receptor alpha variable S-B	XG1079
	zinc finger protein (clone G47)	BE350026
	hypothetical protein FLJ21983	NM_024560
	solute-carrier family 8 (neutrotransmitter transporter, creatinine), member 8	AW276522
	alko-keto reductase family 1, member C4 (chlorocone reductase, 3-alpha hydroxysteroid dehydrogenase, type I; dihydroadiol dehydrogenase 4)	AB045929
	hypothetical protein FLJ11637	NM_024967
	proteinase 3 (serine proteinase, neutrophil, Wegener granulomatosis autoantigen)	NM_002777
	Consensus includes gb:BEB88744 /FEA=EST /DB_XREF=ggi:10345354 /DB_XREF=est:BD151365BF1 /CLONE=IMAGE:3915133 /UG=Hs.169274 ESTs, Highly similar to IF BEB88744	IF BEB88744
	Consensus includes gb:AV0708130 /FEA=EST /DB_XREF=ggi:10725395 /DB_XREF=est:AV0708130 /CLONE=ADCAIA05 /UG=Hs.283965 Homo sapiens scavenger recept AV0708130	AV0708130
	Consensus includes gb:AO333403 /DEF=Human DNA sequence from clone 8EDY on chromosome Xq25-26.3 Contains F9 (coagulation factor IX (plasma thromboplastin co AO333403	AO333403
	VinhAI 547 protein	Al667428
	KIAA1216 protein	NM_004895
	Consensus includes gb:AW663632 /FEA=EST /DB_XREF=ggi:7456210 /DB_XREF=est:h12a03.x1 /CLONE=IMAGE:2981548 /UG=Hs.285825 Homo sapiens mRNA: cDNA AW663632	CNDA AW663632
	telomeric repeat binding factor (NIMA-interacting) 1	A3471136
	chromosome 2 open reading frame 6	AK023321
	HLA-B associated transcript 1	A1282485
	histatin 1	NM_002159
	human epididymis-specific 3 alpha	NM_006663
	emogamnil binding protein (sterol isomerase)	NM_006579
	epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b)) oncogene homolog, avian)	U95089
	prf42 homolog	NM_020640
	Consensus includes gb:AL080233.1 /DEF=Homo sapiens mRNA: cDNA DKFPz586L111 (from clone DKFZP586L111), /FEA=mRNA /DB_XREF=ggi:5262726 /UG=Hs.32656 AL080233	AL080233
	KIAA1216 protein	A017382
	H4 histone family, member A	NM_003538
	Consensus includes gb:BF511231 /FEA=EST /DB_XREF=ggi:11594529 /DB_XREF=est:LJI-H-B4-aqi-g-11-Q.U.I.s1 /CLONE=IMAGE:3085244 /UG=Hs.288582 ESTs, Weakly BF511231	BF511231
	spectrin, beta, erythrocytic (includes spherocytosis, clinical type 1)	NM_000347
	splicing factor, arginine/serine-rich 3	AF107405
	uronyl 2-sulfotransferase	AAW18882
	vinculin	NM_014000
	platelet-derived growth factor receptor, alpha polypeptide	M22734
	fusion, derived from t(12;16) malignant liposarcoma	AW511410
	Consensus includes gb:AKO21983 /DEF=Homo sapiens cDNA FLJ11928 fs, clone HEMBB1000374, /FEA=mRNA /DB_XREF=ggi:10433297 /UG=Hs.306617 Homo sapie AKO21983	AKO21983
	enkephalinase 2, (gamma, neuronal)	NM_001975
	Consensus includes gb:BF674749 /FEA=EST /DB_XREF=ggi:11948644 /DB_XREF=est:602136893 /CLONE=IMAGE:4273270 /UG=Hs.258800 ESTs BF674749	BF674749
	hypothetical protein FLJ13197	NM_024614

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protein phosphatase 2, regulatory subunit B (B56), alpha isoform
 Consensus includes gb:AK024561.1 /DEF=Homo sapiens cDNA: FLJ20908 fls, clone ADSE00417, /FEA=mRNA /DB_XREF=gi:10436870 /UG=Hs.306689 Homo sapiens c AK024561 1.56

hypothetical protein FLJ21302
 growth hormone receptor
 Consensus includes gb:AJ251844.1 /DEF=Homo sapiens partial mRNA for MOZCBP chimeric transcript type II, /FEA=mRNA /GEN=MOZCBP /DB_XREF AJ251844 1.56

transmembrane protein 8 (five membrane-spanning domains)
 Consensus includes gb:X81636.1 /DEF=H. sapiens clathrin light chain a gene
 Consensus includes gb:X81636.1 /DEF=H. sapiens clathrin light chain a gene 1.56

pregnancy specific beta-1-glycoprotein 1
 HBS1-like (S. cerevisiae)
 lymphocyte antigen 8
 sirutin silent mating type information regulation 2 homolog 2 (S. cerevisiae)
 leukocyte tyrosine kinase
 Consensus includes gb:AL355532 /DEF=Human DNA sequence from clone RP11-487F5 on chromosome 6 Contains the 3 part of the GRIK2 (glutamate receptor, ionotropic) AL355532 1.56

Consensus includes gb:AI083505 /FEA=EST /DB_XREF=gi:5810724 /DB_XREF=est:W48h12.x1 /CLONE=IMAGE:2510855 /UG=Hs.137576 ribosomal protein L34 pseudo AI083505 1.56

hypothetical protein DKFZ782186
 FK506 binding protein 6 (38kD)
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily e, member 1
 Consensus includes gb:AV682438 /FEA=EST /DB_XREF=gi:10284299 /DB_XREF=est:AV682436 /CLONE=GKABAE08 /UG=Hs.5848 Homo sapiens mRNA; cDNA DKFZ: AV682436 1.55

Consensus includes gb:AV682438 /FEA=EST /DB_XREF=gi:10284299 /DB_XREF=est:AV682436 /CLONE=GKABAE08 /UG=Hs.5848 Homo sapiens mRNA; cDNA DKFZ: AV682436 1.55

zinc finger protein 358
 Consensus includes gb:AK024158.1 /DEF=Homo sapiens cDNA FLJ14096 fls, clone MAMMA1000752, /FEA=mRNA /DB_XREF=gi:10436469 /UG=Hs.303289 Homo sapiens AK024158 1.55

endothelin 1
 general transcription factor IIIH, polypeptide 1 (62KD subunit)
 collagen, type I, alpha 1
 p53-regulated apoptosis-inducing protein 1
 son of sevenless (Drosophila) homolog 2
 Consensus includes gb:W85912 /FEA=EST /DB_XREF=gi:1398341 /DB_XREF=est:zh52b12.s1 /CLONE=IMAGE:415679 /UG=Hs.323470 Homo sapiens cDNA: FLJ22106 W85912 1.55

wingless-type MMTV Integration site family, member 78
 nuclear fragile X mental retardation protein interacting protein 1
 hypothetical protein DKFZ78181823
 hypothetical protein FLJ22490
 Consensus includes gb:AV717062 /FEA=EST /DB_XREF=gi:10814214 /DB_XREF=est:AV717062 /CLONE=DCBCEG08 /UG=Hs.274553 Homo sapiens cDNA FLJ10906 fl AV717062 1.55

fibroblast growth factor 20
 Consensus includes gb:AL138139 /DEF=Human DNA sequence from clone RP4-78112 on chromosome 6 Contains 3 part of the gene for enhancer of filamentation (HEF1), AL138139 1.55

hypothetical protein FLJ12284
 Consensus includes gb:AK023843.1 /DEF=Homo sapiens cDNA FLJ13781 fls, clone PLACE4000465, /FEA=mRNA /DB_XREF=gi:10435902 /UG=Hs.252820 Homo sapiens AK023843 1.55

retinoblastoma (X-linked, juvenile) 1
 Consensus includes gb:X16454 /DEF=Human gene for carcinoembryonic antigen subdomains A and B /FEA=CDS /DB_XREF=gi:29862 /UG=Hs.248025 Human gene for c X16454 1.55

gb:BC005233.1 /DEF=Homo sapiens, clone MGC:12257, mRNA, complete cds, /FEA=mRNA /PROD=Unknown (protein for MGC:12257) /DB_XREF=gi:13528868 /UG=Hs. BC005233 1.55

myotubularin related protein 1
 hypothetical protein FLJ20874
 fibulin 1 (Marfan syndrome)
 solute carrier family 20 (phosphate transporter), member 1
 Glutamate receptor interacting protein
 kinesin family member 1C
 zinc finger protein 257
 nictalopin
 Consensus includes gb:AK024615.1 /DEF=Homo sapiens cDNA: FLJ20962 fls, clone ADSH00804, /FEA=mRNA /DB_XREF=gi:10436931 /UG=Hs.306698 Homo sapiens c AK024615 1.55

Cdc42 effector protein 4; binder of Rho GTPases 4
 Ewing sarcoma breakpoint region 1
 hypothetical protein DKFZ761N09121
 glutathione peroxidase 2 (gastrintestinal)
 guanine nucleotide binding protein 13, gamma
 cathepsin B
 immunoglobulin lambda locus
 Consensus includes gb:AV652031 /FEA=EST /DB_XREF=gi:3873045 /DB_XREF=est:AV652031 /CLONE=GLCCWA12 /UG=Hs.282344 Homo sapiens cDNA FLJ13387 fl AV652031 1.55

hypothetical protein FLJ10139
 chromosome 15 open reading frame 5
 butyrophilin, subfamily 3, member A2
 UDP glycosyltransferase 1 family, polypeptide A8
 chromosome 11 open reading frame 8
 p300/CBP-associated factor
 AV727449 1.55

a disintegrin and metalloproteinase domain 10
 lefty homolog-like (mouse)
 gb:BC005220.1 /DEF=Homo sapiens, Similar to chaperonin containing TCP1, subunit 8 (theta), clone MGC:12240, mRNA, complete cds. /FEA=mRNA /PROD=Similar to c
 borden
 PRO1621 protein
 Rho GDP dissociation inhibitor (GDI) gamma
 CG1-143 protein
 CD24 antigen (small cell lung carcinoma cluster 4 antigen)
 Consensus includes gb:AK024958.1 /DEF=Homo sapiens cDNA: FLJ21305 fts, clone COL02124. /FEA=mRNA /DB_XREF=gi:10437382 /UG=Hs.287658 Homo sapiens c
 x69397
 hypothetical protein MGC11138
 nerve growth factor receptor (TNFR superfamily, member 16)
 complement component 1, r subcomponent
 synaptotagmin 1
 STAT induced STAT inhibitor 3
 adenosine kinase
 poliovirus receptor
 DKFZP564C1940 protein
 potassium inwardly-rectifying channel, subfamily J, member 9
 TRF-proximal protein
 calpain 5
 3'(2', 5'-bisphosphate nucleotidase 1
 HEF like Protein
 hairy/enhancer-of-split related with YRPW motif 1
 adenovirus 5 E1A binding protein
 DKFZP564O092 protein
 MHC class I polypeptide-related sequence B
 olfactory receptor, family 12, subfamily D, member 2
 RNA binding motif protein 4
 Consensus includes gb:AA443771 /FEA=EST /DB_XREF=gi:2156446 /DB_XREF=est:zw9508.s1 /CLONE=IMAGE:784743 /UG=Hs.270138 ESTs
 KIAA0410 gene product
 cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)
 hypothetical protein FLJ11181
 stoned B/TfIIA-alpha/beta-like factor
 centromere protein C 1
 Consensus includes gb:AK021495.1 /DEF=Homo sapiens cDNA FLJ11433 fts, clone HEMBA1001121. /FEA=mRNA /DB_XREF=gi:10432690 /UG=Hs.287415 Homo sapie
 poliovirus receptor-related 2 (herpesvirus entry mediator B)
 KIAA0798 gene product
 transcription elongation factor (SII) elongin A2
 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3)
 Consensus includes gb:AK021514.1 /DEF=Homo sapiens cDNA FLJ11452 fts, clone HEMBA1001435. /FEA=mRNA /DB_XREF=gi:10432710 /UG=Hs.148598 Homo sapie
 synaptonemal complex protein 1
 a disintegrin and metalloproteinase domain 22
 Consensus includes gb:AL049252.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564D183 (from clone DKFZp564D193). /FEA=mRNA /DB_XREF=gi:4495993 /UG=Hs.3020 AL049252
 nascent helix loop helix 2
 Consensus includes gb:AL080106.1 /DEF=Homo sapiens mRNA: cDNA DKFZp568O053 (from clone DKFZp568O053). /FEA=mRNA /DB_XREF=gi:5262531 /UG=Hs.3063 AL080106
 Consensus includes gb:AK022120.1 /DEF=Homo sapiens cDNA FLJ12058 fts, clone HEMBB1002092. /FEA=mRNA /DB_XREF=gi:10433445 /UG=Hs.296688 Homo sapie
 integrin-binding sialoprotein (bone sialoprotein, bone sialoprotein II)
 Cdc42 effector protein 3
 tumor protein p73
 lyxyl oxidase-like 1
 gb:L14458.1 /DEF=Human Ig rearranged kappa-chain gene V-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398492 /FL=gb:L14458.1
 sulfate carrier family 7 (cationic amino acid transporter, y+ system), member 8
 glycolipid transfer protein
 CATR tumorigenicity conversion 1
 small inducible cytokine A2 (monocyte chemotactic protein 1)
 hypothetical protein FLJ22332
 transcription factor eta
 hypothetical protein MGC2474
 leucine zipper-like protein
 Consensus includes gb:AF038193.1 /DEF=Homo sapiens clone 23608 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2795913 /UG=Hs.6220 Homo sapiens clone 23608 m
 chemokine (C-C motif) receptor-like 2
 reticulocalbin 2, EF-hand calcium binding domain

DKFZp434P211 protein	NM_014549	1.54
early lymphoid activation protein	L22650	1.54
hypothetical protein PRO2032	BE328496	1.54
vitamin A responsive; cytoskeleton related	NM_006407	1.54
putative purinergic receptor	NM_014499	1.54
potassium voltage-gated channel, KQT-like subfamily, member 1	U89364	1.54
A kinase (PRKA) anchor protein 5	NM_004857	1.54
integrin, beta 1 (fibronectin receptor, beta polypeptide, anigen CD29 includes MDF2, MSK12)	BF663461	1.54
protein A	NM_019858	1.54
Consensus includes gb:AL109671.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 29222. /FEA=mRNA /DB_XREF=gi:5689802 /UG=Hs.1408331 AL109671	AL109671	1.54
hypothetical protein	NM_020153	1.54
CSR1 protein	NM_016240	1.54
meprin A, beta	NM_005925	1.54
gamma-glutamyltransferase 1	J05235	1.54
integrin, beta 1 (fibronectin receptor, beta polypeptide, anigen CD29 includes MDF2, MSK12)	M34189	1.54
hypothetical protein FLJ10512	NM_018121	1.54
fibronectin 1	AJ276395	1.54
apolipoprotein C-II	NM_001645	1.54
M-phase phosphoprotein 10 (U3 small nuclear ribonucleoprotein)	AL545921	1.54
Consensus includes gb:AL136094 /DEF=Human DNA sequence from clone RP3-527B10 on chromosome 6q25.1-25.3 Contains a pseudogene similar to HMG (high mobility group) protein 143	AL136094	1.54
tryptase beta 2	AF099143	1.54
glutamate dehydrogenase 1	NM_005271	1.54
fibronectin leucine rich transmembrane protein 2	AF168676	1.54
immunoglobulin kappa constant	BQ482805	1.54
G protein-coupled receptor 22	NM_005295	1.54
chemokine (C-C motif) receptor 1	NM_001295	1.54
bulbos pemphigoid antigen 1 (230/240KD)	AL098710	1.54
hypothetical protein FLJ12439	NM_023077	1.54
polyadenylation binding protein-interacting protein 1	NM_006451	1.54
nebulin	NM_016365	1.54
adrenergic, alpha-1A-, receptor	D32201	1.54
hypothetical protein FLJ20695	A1817074	1.54
coptine VI (neuronal)	AL050397	1.54
trophoblast glycoprotein	NM_006670	1.54
immunoglobulin superfamily, member 4	AF132811	1.54
activator of CREM in testis	AL132776	1.54
Consensus includes gb:AL157428.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761F152 (from clone DKFZp761F152). /FEA=mRNA /DB_XREF=gi:7018458 /UG=Hs.3064; AL157428	AL157428	1.54
Consensus includes gb:AA100250 /FEA=EST /DB_XREF=gi:1647823 /DB_XREF=IMAGE350003 /UG=Hs.90869 Homo sapiens clones 24622 an AA100250	AA100250	1.54
solute carrier family 26, member 4	NM_000441	1.54
ephrin-A3	AW189015	1.54
sodium channel, voltage-gated, type III, alpha polypeptide	AF225986	1.54
hypothetical protein FLJ22287	NM_025135	1.54
KIAA0446 gene product	AL526243	1.54
arachidonate 15-lipoxygenase	NM_001140	1.54
SH3-domain protein 5 (ponatin)	NM_015385	1.54
DKFZP438L151 protein	AW007573	1.54
KIAA1091 protein	AA348595	1.54
Consensus includes gb:AK022192.1 /DEF=Homo sapiens cDNA FLJ12130 fis, clone MAMMA1000251. /FEA=mRNA /DB_XREF=gi:10433534 /UG=Hs.319567 Homo sapiens	AK022192	1.53
hypothetical protein FLJ20502	NM_017845	1.53
zinc finger protein 236	NM_007345	1.53
synaptopodin	NM_007288	1.53
sodium channel, voltage-gated, type XII, alpha polypeptide	AF150882	1.53
Consensus includes gb:AL117520.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434P2435 (from clone DKFZp434P2435). /FEA=mRNA /DB_XREF=gi:5912036 /UG=Hs.304 AL117520	NM_004886	1.53
amyloid beta (A4) precursor protein-binding, family A, member 3 (X11-like 2)	AF130104	1.53
hypothetical protein DKFZp564K142 similar to implantation-associated protein	NM_001430	1.53
endothelial PAS domain protein 1	AW338761	1.53
insulin-like growth factor binding protein, acid labile subunit	NM_002702	1.53
POU domain, class 6, transcription factor 1	NM_005013	1.53
nucleobindin 2	NM_003058	1.53
solute carrier family 22 (organic cation transporter), member 2	AF228163	1.53
nuclear LIM interactor-interacting factor	NM_016204	1.53
growth differentiation factor 2	BC003528	1.53
gb:BC003528.1 /DEF=Homo sapiens, clone MGC:10782, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:10782) /DB_XREF=gi:13087626 /UG=Hs. BC003528		

Consensus includes gb:AU147295 /FEA=EST /DB_XREF=est:AU147295 /CLONE=MAMMA1000264 /UG=Hs.288575 Homo sapiens cDNA FLJ14 AU147295
 protein phosphatase 1, regulatory (inhibitor) subunit 12A
 interferon, alpha-inducible protein 27
 Consensus includes gb:U81981 /DEF=Homo sapiens amiloride-sensitive epithelial sodium channel alpha subunit alpha ENAC-1 and alpha ENAC-2 (SCNN1A) genes, allent
 SRY (sex determining region Y)-box 11
 very low density lipoprotein receptor
 gamma-aminobutyric acid (GABA) receptor, rho 1
 hypothetical protein FLJ22865
 insulin-like growth factor binding protein 5
 glutamate receptor, ionotropic, AMPA 1
 TAF5 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 100 kD
 deleted in bladder cancer chromosome region candidate 1
 Consensus includes gb:AK026493.1 /DEF=Homo sapiens cDNA: FLJ22840 fls, clone KIAU4708 /FEA=mRNA /DB_XREF=gi:10433366 /UG=Hs.287293 Homo sapiens cDI AK026493
 squalene epoxidase
 hypothetical protein FLJ20041
 Consensus includes gb:AK021842.1 /DEF=Homo sapiens cDNA FLJ11780 fls, clone HEMBA1005931, weakly similar to ZINC FINGER PROTEIN 83. /FEA=mRNA /DB_XF
 Insulin-like growth factor 2, antisense
 hyaluronan synthase 1
 hypothetical protein FLJ11196
 tripartite motif-containing 14
 KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2
 G protein-coupled receptor 3
 polymyositis/scleroderma autoantigen 1 (75kD)
 zinc finger protein 79 (pT7)
 Consensus includes gb:AK000220.1 /DEF=Homo sapiens cDNA FLJ20213 fls, clone COLF1971, highly similar to AB007921 Homo sapiens mRNA for KIAA0452 protein. /F
 hypothetical protein DKFZ564D0482
 Consensus includes gb:A1697803 /FEA=EST /DB_XREF=gi:4985703 /DB_XREF=est:wa17605.x1 /CLONE=IMAGE2341376 /UG=Hs.114983 Homo Sapiens (clone 83B3E
 cytokine receptor-like factor 1
 hypothetical protein FLJ22349
 glutamate receptor, ionotropic, kainate 1
 RNA binding motif protein 5
 elcC homolog 1 (E. coli)
 cartilage acidic protein 1
 hypothetical protein FLJ10813
 thyroid stimulating hormone receptor
 oxidative stress induced like
 ryanodine receptor 1 (skeletal)
 LIM domain only 4
 hypothetical protein MGC3771
 solute carrier family 1 (high affinity aspartate/glutamate transporter), member 6
 multifunctional polypeptide similar to SAICAR synthetase and AIR carboxylase
 HDAC2BP protein
 hypothetical protein from BCR42 region
 Consensus includes gb:AK025180.1 /DEF=Homo sapiens cDNA: FLJ21527 fls, clone COL05961 /FEA=mRNA /DB_XREF=gi:10437643 /UG=Hs.306779 Homo sapiens cC
 RPA-binding trans-activator
 myomesin (M-protein) 2 (165kD)
 annexin A4
 hypothetical protein FLJ10280
 STECQ-like kinase
 Consensus includes gb:AK022006.1 /DEF=Homo sapiens cDNA FLJ11944 fls, clone HEMMB1000684 /FEA=mRNA /DB_XREF=gi:10433318 /UG=Hs.300771 Homo sapie
 polymerase (DNA directed), gamma
 GDNF family receptor alpha 3
 tumor necrosis factor (ligand) superfamily, member 10
 phosphoinositide-3-kinase, catalytic, delta polypeptide
 Fzr1 protein
 neuropeptide Y
 Consensus includes gb:AF052095.1 /DEF=Homo sapiens clone 23911 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360401 /UG=Hs.167344 Homo sapiens clone 23911
 transcription factor 3 (E2A Immunoglobulin enhancer binding factors E12/E47)
 DNA segment, numerous copies, expressed probes (GS1 gene)

KIAA0974 protein
 Consensus includes gb:AL117546.1 /DEF=Homo sapiens mRNA; cDNA DKFZ586C1923 (from clone DKFZ586C1923). /FEA=mRNA /DB_XREF=gi:5912079 /UG=Hs.24
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1
 killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2
 aggrscan 1 (chondroitin sulfate proteoglycan 1, large aggregating proteoglycan, antigen identified by monoclonal antibody A0122)
 cutaneous T-cell lymphoma tumor antigen so70-2
 hypothetical protein
 sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)
 thioaurine S-methyltransferase
 Consensus includes gb:AL117451.1 /DEF=Homo sapiens mRNA; cDNA DKFZ586E2317 (from clone DKFZ586E2317). /FEA=mRNA /DB_XREF=gi:5911901 /UG=Hs.28
 hypothetical protein FLJ20716
 hypothetical protein PRO1446
 protocadherin 1 (cadherin-like 1)
 APG5 autophagy 5-like (S. cerevisiae)
 gb:L14453.1 /DEF=Human Ig rearranged mu-chain gene V-H-D-N-J-region, complete cds. /FEA=mRNA /DB_XREF=gi:398494 /FL=gb:L14453.1
 docking protein 2, 50kD
 cell growth regulatory with EF-hand domain
 LAP (leucine-rich repeats and PDZ) and no PDZ protein
 class-I MHC-restricted T cell associated molecule
 likely homolog of rat kinase D-interacting substance of 220 kDa
 Ig superfamily protein
 phosphodiesterase 1C, calmodulin-dependent (70kD)
 phosphothioesterase-3-kinase, regulatory subunit, polypeptide 2 (p85 beta)
 interferon (alpha, beta and omega) Receptor 2
 Consensus includes gb:AA335066 /FEA=EST /DB_XREF=gi:2279319 /DB_XREF=est:nf84a10.51 /CLONE=IMAGE:926586 /UG=Hs.288553 Homo sapiens, clone IMAGE:3 AA335066
 Consensus includes gb:AL181956.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761A17121 (from clone DKFZp761A17121). /FEA=mRNA /DB_XREF=gi:7328008 /UG=Hs.28
 Consensus includes gb:AK001065.1 /DEF=Homo sapiens cDNA FLJ10203 fit, clone HEMBA1004930, moderately similar to 26S PROTEASOME SUBUNIT S5B. /FEA=ntf AK001065
 chromosome X open reading frame 6
 pleckstrin homology, Sec7 and coiled-coil domains, binding protein
 wingless-type MMTV integration site family, member 8B
 KIAA0944 protein
 Consensus includes gb:A927067 /FEA=EST /DB_XREF=gi:5663031 /DB_XREF=est:wo87f01.x1 /CLONE=IMAGE:2462329 /UG=Hs.289068 Homo sapiens cDNA FLJ119
 vinculin
 lysosomal apyrase-like 1
 glycerol-3-phosphate dehydrogenase 1 (soluble)
 SHC-domain, GRB2-like, endophilin B2
 transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)
 hypothetical protein FLJ14075
 hypothetical protein FLJ20313
 guanine nucleotide binding protein 4
 tetraspan NET-6 protein
 Consensus includes gb:AUI56204 /FEA=EST /DB_XREF=gi:11017725 /DB_XREF=est:AUI56204 /CLONE=PLACE:1002583 /UG=Hs.306651 Homo sapiens cDNA FLJ134
 Consensus includes gb:AV702306 /FEA=EST /DB_XREF=gi:10718636 /DB_XREF=est:AV702306 /CLONE=ADBBD05 /UG=Hs.321211 ESTs, Highly similar to zinc finger 1
 protein tyrosine phosphatase, receptor type, D
 discs, large (Drosophila) homolog-associated protein 1
 retinoic acid induced 17
 pleckstrin homology-like domain, family A, member 3
 Consensus includes gb:D25272.1 /DEF=Homo sapiens mRNA, clone RES4-16. /FEA=mRNA /DB_XREF=gi:464187 /UG=Hs.174225 Homo sapiens mRNA, clone RES4-16
 heterogeneous nuclear ribonucleoprotein L
 homolog of mouse BMP-2 inducible kinase
 dynamin 1
 GDNF family receptor alpha 2
 dopamine receptor D1
 gb:BC005043.1 /DEF=Homo sapiens, Similar to tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain, clone MGC:12787, mRNA,
 multiple inositol polyphosphate histidine phosphatase, 1
 nuclear receptor coactivator 1
 fibroblast growth factor 18
 G protein-coupled receptor 20
 olfactory receptor, family 7, subfamily E, member 2 pseudogene
 malignancy-associated protein
 NADH:ubiquinone oxidoreductase MLRO subunit homolog
 homeo box (H8 family) 1
 basic-helix-loop-helix-PAS protein

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thyroid hormone receptor-associated protein, 240 kDa subunit
 hypothetical protein FLJ22870
 scaffold attachment factor B
 hypothetical protein PRO2176
 zinc finger protein 345
 SH3-domain protein 5 (ponsin)
 PDZ-73 protein
 Smc2-related CBP activator protein
 gb:U79250.1 /DEF=Human glycerol-3-phosphate dehydrogenase mRNA, complete cds. /FEA=mRNA /PROD=glycerol-3-phosphate dehydrogenase /DB_XREF=gi:1806827 /U79250
 KIAA0548 protein
 gb:BC002778.1 /DEF=Homo sapiens. Similar to myosin light chain 2, precursor lymphocyte-specific, clone MCC-3479, mRNA, complete cds. /FEA=mRNA /PROD=Similar
 guanine nucleotide binding protein (G protein), alpha 11 (Gq class)
 Consensus includes gb:AF131777.1 /DEF=Homo sapiens clone 25061 mRNA sequence. /FEA=mRNA /DB_XREF=gi:4408602 /UG=Hs.183475 Homo sapiens clone 25061 AF131777
 A kinase (PRKA) anchor protein 1
 KIAA1105 protein
 ecdonucleoside triphosphate diphosphohydrolase 3
 hypothetical protein FLJ20623
 KIAA0557 protein
 DMC1 dosage suppressor of mek1 homolog, meiosis-specific homologous recombination (yeast)
 LOC89231
 ectodermal dysplasia 1, anhidrotic
 ATP-binding cassette, sub-family B (MDR/TAP), member 11
 ELAV (embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D)
 Consensus includes gb:AA78300 /FEA=EST /DB_XREF=gi:4371526 /DB_XREF=est:tm39e01.x1 /CLONE=IMAGE:2160504 /UG=Hs.192789 ESTs, Weakly similar to ALU AF478300
 hypothetical protein FLJ10803
 purinergic receptor P2Y, G-protein coupled, 2
 likely ortholog of mouse SRY-box containing gene 17
 medium-chain acyl-CoA synthetase
 KIAA0429 gene product
 arsenate resistance protein ARS2
 hypothetical protein FLJ20700
 ATPase, Class V, type 10C
 PRO3053 protein
 Consensus includes gb:AL080086.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J1762 (from clone DKFZp564J1762). /FEA=mRNA /DB_XREF=gi:5262502 /UG=Hs.274 AL080086
 mitogen-activated protein kinase kinase kinase 12
 protein kinase, cGMP-dependent, type 1
 follistatin-like 1
 Consensus includes gb:AL035087 /DEF=Human DNA sequence from clone 170F5 on chromosome Xq22.3-24. Contains an HMGI (high-mobility group (nonhistone) chromatin) domain
 Consensus includes gb:AK024896.1 /DEF=Homo sapiens cDNA: FLJ21243 (fs, clone COL01164). /FEA=mRNA /DB_XREF=gi:10437310 /UG=Hs.268016 Homo sapiens CC AK024896
 cadherin, EGF LAG seven-pass G-type receptor 1 (flamingo homolog, Drosophila)
 hypothetical protein FLJ11583
 solute carrier family 22 (organic cation transporter), member 5
 Consensus includes gb:AL359553 /DEF=Human DNA sequence from clone RPS-871G17 on chromosome 1. Contains ESTs, STSs and GSSs. Contains the HAQ2 gene for AL359553
 Consensus includes gb:AL080112.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586H0722 (from clone DKFZp586H0722). /FEA=mRNA /DB_XREF=gi:5262539 /UG=Hs.33 AL080112
 Consensus includes gb:AK022331.1 /DEF=Homo sapiens cDNA: FLJ12269 (fs, clone MAMMA1001635). /FEA=mRNA /DB_XREF=gi:10433706 /UG=Hs.283925 Homo sapiens AK022331
 Consensus includes gb:U80770.1 /DEF=Human EST clone 251800 mariner transposon Hamar1 sequence. /FEA=mRNA /DB_XREF=gi:2921608 /UG=Hs.42192 Human ES U80770
 KIAA0877 gene product
 ecotropic viral integration site 2B
 low density lipoprotein-related protein 2
 2-5-oligoadenylate synthetase-like
 hypothetical protein FLJ10718
 N-acetylated alpha-linked acidic dipeptidase-like; iLEAL DIPEPTIDYLPEPTIDASE
 aldehyde dehydrogenase 3 family, member B1
 hypothetical protein
 hypothetical protein MGC13053
 Consensus includes gb:AA131826 /FEA=EST /DB_XREF=gi:1693379 /DB_XREF=est:z37f06.s1 /CLONE=IMAGE:504131 /UG=Hs.107164 spectrin, beta, non-erythrocytic
 KIAA0227 protein
 toll-like receptor 2
 solute carrier family 7 (cationic amino acid transporter, y+ system), member 4
 cell division cycle 2-like 2
 CREBBP/EP300 inhibitory protein 1
 RAB2, member RAS oncogene family
 chromosome 3p21.1 gene sequence
 NM_013286

drebrin 1
 hypothetical protein FLJ14009
 hydroxysteroid (17-beta) dehydrogenase 2
 homeo box 11 (T-cell lymphoma 3-associated breakpoint)
 CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 1
 Consensus includes gb:BF591638 /FEA=EST /DB_XREF=gi:11683962 /DB_XREF=est:7n5g03.x1 /CLONE=IMAGE:3568325 /UG=Hs.129663 ESTs
 paternally expressed 3
 Consensus includes gb:AA4770170 /FEA=EST /DB_XREF=gi:2821408 /DB_XREF=est:ah84d09.s1 /CLONE=1322321 /UG=Hs.288156 Homo sapiens cDNA: FLJ21819 fs.
 lental (3) malignant brain tumor (3)mbt protein (Drosophila) homolog
 Consensus includes gb:AK024568.1 /DEF=Homo sapiens cDNA: FLJ20915 fs, done ADSE00892, /FEA=mRNA /DB_XREF=gi:0438878 /UG=Hs.306891 Homo sapiens cDNA: AK024568
 epidermal growth factor (beta-urogastrone)
 protein phosphatase 1, catalytic subunit, beta isoform
 CDC14 cell division cycle 14 homolog B (S. cerevisiae)
 G antigen 7B
 voltage-gated sodium channel beta-3 subunit (scn3b gene)
 uridine monophosphate synthetase (uridate phosphoribosyl transferase and orotidine-5'-decarboxylase)
 KIAA1751 protein
 Consensus includes gb:A1739035 /FEA=EST /DB_XREF=gi:5101018 /DB_XREF=est:w34e04.x1 /CLONE=IMAGE:2392158 /UG=Hs.28974 ESTs
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3
 Consensus includes gb:N34842 /FEA=EST /DB_XREF=gi:1155984 /DB_XREF=est:ty45d11.s1 /CLONE=IMAGE:276501 /UG=Hs.6580 Homo sapiens cDNA: FLJ23227 fs
 hypothetical protein FLJ21919
 lental (3) malignant brain tumor (3)mbt protein (Drosophila) homolog
 hypothetical protein FLJ13059
 Consensus includes gb:AF234255.1 /DEF=Homo sapiens clone KM36 immunoglobulin light chain variable region mRNA, partial cds. /FEA=mRNA /PROD=immunoglobulin
 solute carrier family 10 (sodium/bile acid cotransporter family), member 1
 syntaxin 5A
 Consensus includes gb:W72053 /FEA=EST /DB_XREF=gi:1382323 /DB_XREF=est:zd66e04.s1 /CLONE=IMAGE:345830 /UG=Hs.110771 Homo sapiens cDNA: FLJ21904
 nuclear receptor co-repressor 2
 KIAA0584 protein
 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit e
 similar to S68401 (cattle) glucose induced gene
 follicular lymphoma variant translocation 1
 Consensus includes gb:AL359578.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547N163 (from clone DKFZp547N163). /FEA=mRNA /DB_XREF=gi:8655637 /UG=Hs.3066
 hypothetical protein LOC37821
 Ser-Thr protein kinase related to the myotonic dystrophy protein kinase
 hypothetical protein MGC14378
 ankyrin repeat and SOCS box-containing 4
 Cluster ind. AJ400621:tp49g02.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2112146 /clone_end=3, /gb=AJ400621 /gi=4243708 /ug=Hs.213540 /len=529
 GIOT-2 for gonadotropin inducible transcription repressor-2
 similar to S. pombe dim1+
 Consensus includes gb:BF512531 /FEA=EST /DB_XREF=gi:11597710 /DB_XREF=est:U1-H-BW1-ami-b-05-O-UJ.s1 /CLONE=IMAGE:3069633 /UG=Hs.297765 ESTs
 geneathionin 1
 hypothetical protein FLJ10035
 Consensus includes gb:AL049342.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566A183 (from clone DKFZp566A183). /FEA=mRNA /DB_XREF=gi:4500128 /UG=Hs.2745
 PR domain containing 10
 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsin, antitrypsin), member 1
 nuclear receptor subfamily 4, group A, member 2
 Consensus includes gb:AL117830.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D115 (from clone DKFZp434D115). /FEA=mRNA /DB_XREF=gi:5912213 /UG=Hs.2527
 CGI-88 protein
 pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 4
 hypothetical protein FLJ11348
 KIAA1887 protein
 transient receptor potential channel 6
 lymphocyte antigen 117
 ring finger protein (C3HC3 type) 6
 kinesin family member 3C
 Consensus includes gb:BF111288 /FEA=EST /DB_XREF=gi:10940958 /DB_XREF=est:7174d02.x1 /CLONE=IMAGE:3527090 /UG=Hs.12438 Homo sapiens cDNA: FLJ228 BF111288
 Consensus includes gb:AL117520.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434P2435 (from clone DKFZp434P2435). /FEA=mRNA /DB_XREF=gi:5912036 /UG=Hs.30 AL117520
 cholecystokinin A receptor
 achaete-scute complex-like 1 (Drosophila)
 neuropeptide Y receptor Y2
 Consensus includes gb:AF023203.1 /DEF=Homo sapiens homeobox protein Ogl 12 (OGL12) mRNA, partial cds. /FEA=mRNA /GEN=OGL12 /PROD=homeobox protein Ogl
 poliovirus receptor
 Consensus includes gb:AF023203.1 /DEF=Homo sapiens homeobox protein Ogl 12 (OGL12) mRNA, partial cds. /FEA=mRNA /GEN=OGL12 /PROD=homeobox protein Ogl

uroplakin 1B	NM_006952	1.50
LAT1-3TM protein	AC003007	1.50
synaptotagmin V	NM_003180	1.50
period homolog 3 (Drosophila)	Z98884	1.50
tumor necrosis factor (ligand) superfamily, member 9	NM_003811	1.50
hypothetical protein FLJ22269	AK025922	1.50
solute carrier family 12 (potassium/chloride transporters), member 4	A017690	1.50
Fzr1 protein	A173260	1.50
VAMP (vesicle-associated membrane protein)-associated protein B and C	NM_004738	1.50
Cluster incl. A1557319:PT2.1, 16, F08.1 Homo sapiens cDNA, 3' end / clone_end=3 / (gb=A1557319 / gi=4489682 / ug=Hs.21921 / len=876	A1557319	1.50
Consensus includes gb:AL080068.1 / DEF=Homo sapiens mRNA, cDNA DKFZp564J062 (from clone DKFZp564J062), / FEA=mRNA / DB_XREF=gi:5282475 / UG=Hs.27253	AL080068	1.50
apolipoprotein B (including Ag(x) antigen)	NM_000384	1.50
chimerin (chimerin) 2	L29126	1.50
SEC24 related gene family, member A (S. cerevisiae)	AJ131244	1.50
glycine amidinotransferase (L-arginine:glycine amidinotransferase)	NM_001482	1.50
autism-related protein 1	AK025298	1.50
adenosine monophosphate deaminase 2 (isoform L)	A016249	1.50
hypothetical protein FLJ23231	NM_025079	1.50
Consensus includes gb:AL137292.1 / DEF=Homo sapiens mRNA, cDNA DKFZp434L1016 (from clone DKFZp434L1016), / FEA=mRNA / DB_XREF=gi:6807752 / UG=Hs.274	AL137292	1.50
Consensus includes gb:AK022322.1 / DEF=Homo sapiens cDNA FLJ12260 fs. clone MAMMA100155.1 / FEA=mRNA / DB_XREF=gi:10433693 / UG=Hs.306634	Homo sapiens AK022322	1.50
siatrophin (gpl.115, leukostallin, CD43)	X69502	1.50
gaslin-releasing peptide	NM_002091	1.50
hypothetical protein DKFZp547H025	NM_020181	1.50
zinc finger protein 222	NM_013360	1.50
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsin, antitrypsin), member 5	J02639	1.50
RYK receptor-like tyrosine kinase	NM_002956	1.50
myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))	U55209	1.50
PRP4/STKWD splicing factor	A118402	1.50
prothymosin, alpha (gene sequence 28)	BF686442	1.50
hypothetical protein FLJ23047	NM_024548	1.50
sulfotransferase family, cytosolic, 2B, member 1	NM_004605	1.50
diaphanous homolog 2 (Drosophila)	NM_007309	1.50
Consensus includes gb:BE892698 / FEA=EST / DB_XREF=gi:10353302 / DB_XREF=est6014339 / TF1 / CLONE=IMAGE:3918989 / UG=Hs.129709	Homo sapiens chromosome BE892698	1.50
Consensus includes gb:N30878 / FEA=EST / DB_XREF=gi:1149398 / DB_XREF=est:1149398 / CLONE=IMAGE:238129 / UG=Hs.82276	trichohyalin N30878	1.50
flotillin 1	AA507012	1.50
platelet-derived growth factor receptor, beta polypeptide	NM_002609	1.50
coagulation factor II (thrombin)	NM_000506	1.50
ankyrin 3, node of Ranvier (ankyrin G)	NM_001149	1.50
tetraspan 5	AA748177	1.50
Consensus includes gb:A1825798 / FEA=EST / DB_XREF=gi:54464689 / DB_XREF=est:18e08.x1 / CLONE=IMAGE:2076038 / UG=Hs.211388	Homo sapiens BAC clone CTE A1825798	1.50
chromosome 11 open reading frame 9	AC004770	1.50
solute carrier family 22 (organic anion transporter), member 7	AF210455	1.50
pp21 homolog	NM_016303	1.50
hypothetical protein FLJ10261	NM_018043	1.50
oligodactyl receptor, family 3, subfamily A, member 2	NM_002551	1.49
FXD domain-containing ion transport regulator 2	NM_001680	1.49
TSPY-like	AL136629	1.49
Consensus includes gb:AL359563.1 / DEF=Homo sapiens mRNA, cDNA DKFZp434D0326 (from clone DKFZp434D0326), / FEA=mRNA / DB_XREF=gi:8655619 / UG=Hs.30	AL359563	1.49
PTD011 protein	NM_014051	1.49
Consensus includes gb:AL137606.1 / DEF=Homo sapiens mRNA, cDNA DKFZp434H0211 (from clone DKFZp434H0211), / FEA=mRNA / DB_XREF=gi:6808354 / UG=Hs.15	AL137606	1.49
flap structure-specific endonuclease 1	BC000323	1.49
glucagon-like peptide 2 receptor	NM_004246	1.49
old astrocyte specifically induced substance	BC328407	1.49
promyelocytic leukemia	AF230410	1.49
glutamate-ammmonia ligase (glutamine synthase)	U08626	1.49
H3 histone family, member K	NM_003538	1.49
KIAA0881 protein	AB023198	1.49
erythrocyte membrane protein band 4.1-like 1	AA573523	1.49
major histocompatibility complex, class II, DQ beta 1	M24364	1.49
gb:AF-098641.1 / DEF=Homo sapiens CD44 isoform RC (CD44) mRNA, complete cds. / FEA=mRNA / GEN=CD44 / PROD=CD44 isoform RC / DB_XREF=gi:3832517 / UG=Hs.1	AF098641	1.49
sin3u1 silent mating type information regulation 2 homolog 4 (S. cerevisiae)	NM_012240	1.49
Consensus includes gb:U05589.1 / DEF=Human ribosomal protein S1 homolog mRNA, partial cds. / FEA=mRNA / DB_XREF=gi:497001 / UG=Hs.371	ribosomal protein S1-III U05589	1.49
myristoylated alanine-rich protein kinase C substrate	AA770596	1.49

calcitonin receptor
desmoplakin (DPI, DPII)
hypothetical protein MGC3178
thymase 1, mast cell
protein phosphatase 1, regulatory (inhibitor) subunit 1A
fibroblast activation protein, alpha
Consensus includes gb:AK000122.1 /DEF=Homo sapiens cDNA FLJ20115 fs, clone COL05594, /FEA=mRNA /DB_XREF=gi:7020002 /UG=Hs.306382 Homo sapiens cDNA AK000122
T-box 2
hypothetical protein FLJ21280
Consensus includes gb:BF511164 /FEA=EST /DB_XREF=gi:11594462 /DB_XREF=est:U1-H-B14-aoi-a-09-0-U1.s1 /CLONE=IMAGE:3084952 /UG=Hs.293309 ESTs
transmembrane 6 superfamily member 2
KIA0953 protein
macrophage stimulating 1 receptor (c-met-related tyrosine kinase)
erythrocyte protein band 4.1-like 4
Consensus includes gb:X83749 /DEF=H.sapiens DNA for Rod transducin alpha-subunit /FEA=mRNA /DB_XREF=gi:37049 /UG=Hs.51147 guanine nucleotide binding prote X63749
S100 calcium binding protein A8 (calgranulin A)
hypothetical protein FLJ20043
protein phosphatase 3 (formerly 2B), catalytic subunit, alpha isoform (calcineurin A alpha)
keratin 7
mitochondrial ribosomal protein L9
paired mesoderm homeo box 1
hypothetical protein MGC3136
ABO blood group (transferase A, alpha 1-3-N-acetylgalactosaminyltransferase; transferase B, alpha 1-3-galactosyltransferase)
TATA box binding protein (TBP)-associated factor, RNA polymerase I, B, 63KD
NK6 transcription factor homolog A (Drosophila)
Consensus includes gb:A824078 /FEA=EST /DB_XREF=gi:5444749 /DB_XREF=est:w35h06.x1 /CLONE=IMAGE:2404859 /UG=Hs.295448 ESTs, Moderately similar to B
H3 histone family, member J
milk fat globule-EGF factor 8 protein
peroxisomal biogenesis factor 11A
ectoderm microtubule-associated protein-like
adrenergic, alpha-1A, receptor
peptidylglycine alpha-amidating monooxygenase
centaurin, delta 1
caldesmon 1
Consensus includes gb:AL353942.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761L23121 (from clone DKFZp761L23121) /FEA=mRNA /DB_XREF=gi:7669981 /UG=Hs.3
hypothetical protein FLJ11856
sulfolipase family, cytosolic, 1C, member 1
phosphate cytidylyltransferase 1, choline, beta isoform
immunoglobulin kappa constant
POU domain, class 4, transcription factor 3
methyl CpG binding protein 2 (Rett syndrome)
prostate androgen-regulated transcript 1
cytochrome P450, subfamily IIC (methoxytoxin 4-hydroxylase), polypeptide 9
interleukin 10 receptor, alpha
KIA0553 protein
Consensus includes gb:A054381 /FEA=EST /DB_XREF=gi:3322168 /DB_XREF=est:q8409.x1 /CLONE=IMAGE:1861265 /UG=Hs.176139 ESTs
Consensus includes gb:AA830143 /FEA=EST /DB_XREF=gi:2903242 /DB_XREF=est:oc44e11.s1 /CLONE=IMAGE:1352588 /UG=Hs.15535 Homo sapiens clone 24582 m
COP9 homolog
Consensus includes gb:AA830143 /FEA=EST /DB_XREF=gi:2903242 /DB_XREF=est:oc44e11.s1 /CLONE=IMAGE:1352588 /UG=Hs.15535 Homo sapiens clone 24582 m
calponin 1, basic, smooth muscle
Consensus includes gb:AU147402 /FEA=EST /DB_XREF=gi:11008923 /DB_XREF=est:AU147402 /CLONE=MAMMA1000567 /UG=Hs.287496 Homo sapiens cDNA FLJ12 AU147402
extracellular glycoprotein EMILIN-2 precursor
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 1
matrix metalloproteinase 7 (matrilysin, uterin)
LIS1-interacting protein NUDE1, rat homolog
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 6 (RNA helicase, 54KD)
Cip1-interacting zinc finger protein
hypothetical protein FLJ12748
ORF
hypothetical protein MGC12488
signal transducer and activator of transcription 2, 113KD
Consensus includes gb:AL050391.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586A181 (from clone DKFZp586A181); partial cds. /FEA=mRNA /GEN=DKFZp586A181 /PI AL050391
gb:UA8722.1 /DEF=Human epidermal growth factor receptor precursor (EGFR) mRNA, complete cds. /FEA=mRNA /PROD=epidermal growth factor receptor precursor /DE UA8722

Consensus includes gb:AL034450 /DEF=Human DNA sequence from clone 11SK14 on chromosome Xq22.3-23 Contains high mobility group protein 2a, ESTs, STS /FEA= AL034450
 Mitochondrial Acyl-CoA Thioesterase
 G protein-coupled receptor kinase 6
 Consensus includes gb:AK022235.1 /DEF=Homo sapiens cDNA FLJ12173 fls, clone MAMMA1000696, /FEA=mRNA /DB_XREF=gi:10433588 /UG=Hs.306630 Homo sapiens
 chondroitin sulfate proteoglycan 2 (versican)
 angiotensin 2
 Consensus includes gb:BE858194 /FEA=EST /DB_XREF=gi:10372818 /DB_XREF=est:7g19c05.x1 /CLONE=IMAGE:3306920 /UG=Hs.323053 Homo sapiens mRNA full le
 BE858194
 U55208
 Consensus includes gb:BE858194 /FEA=EST /DB_XREF=gi:10372818 /DB_XREF=est:7g19c05.x1 /CLONE=IMAGE:3306920 /UG=Hs.323053 Homo sapiens mRNA full le
 BE858194
 U55208
 myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))
 thymosin, beta, identified in neuroblastoma cells
 DnaJ (Hsp40) homolog, subfamily B, member 5
 small inducible cytokine subfamily C, member 2
 hypothetical protein MGC5338
 FE65-LIKE 2
 hypothetical protein FLJ10707
 fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson
 M80634
 histamine receptor H1
 cylindromatosis (turban tumor syndrome)
 membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2)
 hypothetical protein MGC11266
 hypothetical protein FLJ20006
 Leman coiled-coil protein
 cAMP responsive element binding protein-like 1
 ring finger protein 3
 G2 protein
 hyaluronoglucosaminidase 2
 CD18 antigen, b polypeptide
 hypothetical protein FLJ12800
 G antigen 7
 caspase recruitment domain family, member 9
 epithelial V-like antigen 1
 nuclear receptor subfamily 0, group B, member 2
 c-myc binding protein
 G protein-coupled receptor kinase 7
 Fas (TNFRSF6)-associated via death domain
 calpain 3, (p94)
 Consensus includes gb:AL033538 /DEF=Human DNA sequence from clone RP3-477H23 on chromosome 22q12.1-12.2 Contains parts of one or two novel genes, ESTs, S
 AL033538
 Consensus includes gb:AI911273 /FEA=EST /DB_XREF=gi:5631009 /DB_XREF=est:wg3309.x1 /CLONE=IMAGE:2368921 /UG=Hs.122158 ESTs
 lipoma HMGIC fusion partner
 hypothetical protein FLJ21276
 protocadherin alpha 12
 Consensus includes gb:AK021457.1 /DEF=Homo sapiens cDNA FLJ11395 fls, clone HEMBA1000594, /FEA=mRNA /DB_XREF=gi:10432649 /UG=Hs.306595 Homo sapiens
 AK021457
 ubiquitin-like 1 (sentrin)
 apoptotic protease activating factor
 kallikrein 13
 KIAA0589 protein
 CGI-42 protein
 protein tyrosine phosphatase, non-receptor type substrate 1
 erythrocyte membrane protein band 4.1-like 2
 runt-related transcription factor 2
 zinc finger protein 144 (Met-18)
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 24
 clone FLB1727
 E3 ubiquitin ligase SMURF1
 potassium inwardly-rectifying channel, subfamily J, member 5
 surfactant, pulmonary-associated protein B
 chromosome 17 open reading frame 1A
 proliferation-associated ZC4, 38kD
 steroid-5-alpha-reductase, alpha polypeptide 2 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 2)
 epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)
 apolipoprotein L, 2
 isolated-like kinase 1
 potassium voltage-gated channel, Shal-related subfamily, member 2
 Consensus includes gb:BF592058 /FEA=EST /DB_XREF=gi:11684382 /DB_XREF=est:7n9g09.x1 /CLONE=IMAGE:3572920 /UG=Hs.86405 Homo sapiens mRNA, cDNA BF592058

fascin (Strongylocentrotus purpuratus) homolog 3 (actin-bundling protein, testicular)
 glial cell derived neurotrophic factor
 glutathione peroxidase 3 (plasma)
 MAD3 box transcription enhancer factor 2, polypeptide D (myocyte enhancer factor 2D)
 laminin, alpha 4
 hypothetical protein MGC3101
 ubiquitin specific protease 6 (Tre-2 oncogene)
 Consensus includes gb:A116064 /FEA=EST /DB_XREF=gi:3736702 /DB_XREF=est:q650h05.x1 /CLONE=IMAGE:1742457 /UG=Hs.170212 ESTs, Highly similar to b-TRC A116064
 cysteine and glycine-rich protein 2
 amine oxidase, copper containing 3 (vascular adhesion protein 1)
 ubiquitin associated and SH3 domain containing, A
 putative dimethyladenosine transferase
 Consensus includes gb:U52428 /FEA=Human fatty acid synthase gene, partial cds /FEA=CDS_2 /DB_XREF=gi:1374810 /UG=Hs.83190 fatty acid synthase
 T-cell receptor interacting molecule
 dopamine receptor D3
 protein kinase C-like 2
 chromosome 1, open reading frame 9
 bone morphogenetic protein 8 (osteogenic protein 2)
 Consensus includes gb:B166134 /FEA=EST /DB_XREF=gi:12335349 /DB_XREF=est:80228912.F1 /CLONE=IMAGE:4357349 /UG=Hs.250723 FK506 binding protein 12 BF968134
 collagen, type VII, alpha 1 (epidermolysis bullosa, dystrophic, dominant and recessive)
 besonudin
 hypothetical protein FLJ14117
 Consensus includes gb:K00627.1 /DEF=human knoi repeat mma (cdna clone pcd-kpnt-8), 3 end. /FEA=mRNA /DB_XREF=gi:337653 /UG=Hs.203776 Human knoi repeat r K00627
 CD24 antigen (small cell lung carcinoma cluster 4 antigen)
 P311 protein
 Consensus includes gb:A800464.1 /DEF=Homo sapiens mRNA, exon 1, 2, 3, 4, clone:RES4-24A. /FEA=mRNA /DB_XREF=gi:1843395 /UG=Hs.104258 Homo sapiens m A800464
 Consensus includes gb:AK026820.1 /DEF=Homo sapiens cDNA: FLJ23187 fls, clone LNG09902. /FEA=mRNA /DB_XREF=gi:10439765 /UG=Hs.287730 Homo sapiens cC AK026820
 glutamate receptor, ionotropic, N-methyl D-aspartate 2B
 mitogen-activated protein kinase kinase 8
 KIAA1025 protein
 gb:AF130057.1 /DEF=Homo sapiens clone FLB5539 PRO1454 mRNA, complete cds. /FEA=mRNA /PROD=PRO1454 /DB_XREF=gi:11493420 /UG=Hs.323288 Homo sap AF130057
 Consensus includes gb:A1143919 /FEA=EST /DB_XREF=gi:3665728 /DB_XREF=est:q6912.x1 /CLONE=IMAGE:737647 /UG=Hs.193783 Human DNA sequence from c A1143919
 Consensus includes gb:AK001127.1 /DEF=Homo sapiens cDNA FLJ10265 fls, clone HEMBB1001014. /FEA=mRNA /DB_XREF=gi:7022192 /UG=Hs.253369 Homo sapient AK001127
 Consensus includes gb:A480014 /FEA=EST /DB_XREF=gi:4373182 /DB_XREF=est:tm71609.x1 /CLONE=IMAGE:2163945 /UG=Hs.13415 Homo sapiens clone 24571 mf A480014
 modulator of apoptosis 1
 Consensus includes gb:N21364 /FEA=EST /DB_XREF=gi:1126534 /DB_XREF=est:ty50h10.s1 /CLONE=IMAGE:265219 /UG=Hs.26968 Homo sapiens mRNA from chrone N21364
 offactory receptor, family 3, subfamily A, member 3
 dystrubrevin, beta
 thymosin, beta 4, Y chromosome
 transmembrane phosphatase with tensin homology
 colony stimulating factor 3 (granulocyte)
 Consensus includes gb:AW188214 /FEA=EST /DB_XREF=gi:6462650 /DB_XREF=est:xp93g09.x1 /CLONE=IMAGE:2664832 /UG=Hs.250389 ESTs, Weakly similar to AS8: AW188214
 adenomatous polyposis coli like
 aspartyl-IRNA synthetase
 pipercolic acid oxidase
 hypothetical protein FLJ11286
 KIAA0475 gene product
 Consensus includes gb:U82306.1 /DEF=Homo sapiens unknown protein mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1938333 /UG=Hs.123081 Homo U82306
 UV-B repressed sequence, HUR 7
 Consensus includes gb:AW156899 /FEA=EST /DB_XREF=gi:6228300 /DB_XREF=est:tau88c11.x1 /CLONE=IMAGE:2783444 /UG=Hs.12520 Homo sapiens clone 23568, ; AW156899
 ubiquitin-specific processing protease
 bromodomain adjacent to zinc finger domain, 2A
 H3 histone family, member D
 Consensus includes gb:AV756729 /FEA=EST /DB_XREF=gi:10914577 /DB_XREF=est:AV756729 /CLONE=BMCACB11 /UG=Hs.317661 ESTs
 disrupter of silencing 10
 CCR4-NOT transcription complex, subunit 2
 matrix metalloproteinase 14 (membrane-inserted)
 similar to S. cerevisiae RER1
 glycoprotein VI (platelet)
 ribosomal protein L17
 Consensus includes gb:AK022303.1 /DEF=Homo sapiens cDNA FLJ12241 fls, clone MAMMA1001274. /FEA=mRNA /DB_XREF=gi:10433670 /UG=Hs.287503 Homo sapie AK022303
 signal transducer and activator of transcription 4
 transition protein 1 (during histone to protamine replacement)
 NM_003284

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chromosome 4 open reading frame 6	NM_005750	1.47
PCTAIRE protein kinase 1	BE787860	1.47
down-regulator of transcription 1, TBP-binding (negative cofactor 2)	AW516932	1.47
hypothetical protein FLJ20307	NM_017757	1.47
phosphotriesterase related	NM_030664	1.47
interleukin 13 receptor, alpha 2	NM_000640	1.47
microtubule-associated protein tau	J03778	1.47
twist homolog (acrocephalosyndactyl 3; Saethre-Chotzen syndrome) (Drosophila)	X89268	1.47
U2 small nuclear ribonucleoprotein auxiliary factor (65KD)	NM_007279	1.47
AD-O12 protein	NM_016449	1.47
coagulation factor VII (serum prothrombin conversion accelerator)	NM_000131	1.47
insulinoma-associated 1	NM_002196	1.47
masitinib-like 1 (Drosophila)	NM_014757	1.47
a disintegrin and metalloproteinase domain 3a (cyrilstin 1)	X89657	1.47
hypothetical protein FLJ11088	NM_018318	1.47
Consensus includes gb:AK026045.1 /DEF=Homo sapiens cDNA: FLJ22392 fls, clone HRC07868. /FEA=mRNA /DB_XREF=gi:10438761 /UG=Hs.275464 Homo sapiens cDNA: AK026045	AF112372	1.47
TJ6 protein	NM_018519	1.47
Consensus includes gb:AL157398 /DEF=Human DNA sequence from clone RP11-56H7 on chromosome 10. Contains ESTs, STSs and GSSs. Contains the gene for the ne	AL157398	1.47
hypothetical protein PRO2286	AK026097	1.47
hypothetical protein FLJ20013	AI752257	1.47
zinc finger protein 38 (KOX 25)	AU145019	1.47
KIAA1013 protein	U86145	1.47
CD6 antigen	NM_012404	1.47
pp32 related 2	AL047879	1.47
Consensus includes gb:AL047879 /FEA=EST /DB_XREF=gi:4728067 /DB_XREF=est:DKFZp586N1222 /UG=Hs.194251 ESTs, Weakly simi	AL047879	1.47
hypothetical protein FLJ11112	NM_018327	1.47
sepin 6	AL568374	1.47
DnaJ (Hsp40) homolog, subfamily C, member 6	NM_014787	1.47
high-mobility group (nucleosome chromosomal) protein 17-like 3	BC001282	1.47
pancreas-enriched phospholipase C	NM_016341	1.47
zinc finger protein 253	NM_021047	1.47
checkpoint suppressor 1	NM_005197	1.47
lysyl oxidase	L16895	1.47
variable charge, Y chromosome, 2	NM_004678	1.47
DnaJ (Hsp40) homolog, subfamily C, member 4	AW024467	1.47
CD6 antigen	NM_008725	1.47
solute carrier family 34 (sodium phosphate), member 1	NM_003052	1.47
cdg1 protein	NM_014477	1.47
ankyrin repeat domain 6	BE677131	1.46
Ets2 repressor factor	NM_006494	1.46
homeo box D3	Y09880	1.46
Consensus includes gb:AA493300 /FEA=EST /DB_XREF=gi:2223141 /DB_XREF=est:ng97e11.s1 /CLONE=IMAGE:942764 /UG=Hs.104258 Homo sapiens mRNA, exon 1	AA493300	1.46
G antigen 1	NM_001468	1.46
hypothetical protein	NM_012066	1.46
angiotensin receptor 1	NM_004835	1.46
membrane-bound transcription factor protease, site 1	NM_003781	1.46
killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4	AF002256	1.46
asparry-IRNA synthetase	BC000829	1.46
hypothetical protein FLJ21075	NM_025031	1.46
secreted frizzled-related protein 4	AV089415	1.46
prothymosin, alpha (gene sequence 28)	AF348514	1.46
caspase 10, apoptosis-related cysteine protease	NM_001230	1.46
CDC14 cell division cycle 14 homolog B (S. cerevisiae)	AF064105	1.46
transcriptional co-activator with PDZ-binding motif (TAZ)	AA081084	1.46
Consensus includes gb:BC005092.1 /DEF=Homo sapiens, clone IMAGE:3843012, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3843012) /DB_XI	BC005092	1.46
SEC10-like 1 (S. cerevisiae)	NM_006544	1.46
Consensus includes gb:AW204383 /FEA=EST /DB_XREF=gi:6503855 /DB_XREF=est:UH-B1-adv-c-03-0-U1.s1 /CLONE=IMAGE:2717957 /UG=Hs.312723 ESTs, Highly	AW204383	1.46
glucokinase (hexokinase 4, maturity onset diabetes of the young 2)	M69051	1.46
Consensus includes gb:AL098700 /DEF=Human DNA sequence from clone RP3-499B10 on chromosome Xp22.31-22.13. Contains the 3 end of the gene for serpinthreontir	AL098700	1.46
hypothetical protein FLJ10116	NM_018000	1.46
zinc finger protein SBZF3	NM_020394	1.46
Epstein-Barr virus induced gene 2 (lymphocyte-specific G protein-coupled receptor)	NM_004951	1.46
Consensus includes gb:AK000786.1 /DEF=Homo sapiens cDNA FLJ20779 fls, clone COL05077. /FEA=mRNA /DB_XREF=gi:7021088 /UG=Hs.321583 Homo sapiens cDNA AK000786	AK000786	1.46

Consensus includes gb:AL080130.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434E033 (from clone DKFZp434E033). /FEA=mRNA /DB_XREF=gi:5262569 /UG=Hs.1574/AL080130 1.46
RNA binding motif, single stranded interacting protein 2 AAS36000 1.46
KIAA1473 protein AB040906 1.46
gb:BC001217.1 /DEF=Homo sapiens. Similar to KIAA0029 protein, clone MSG:767, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0029 protein /DB_XREF=g 1.46
AL136883 1.46
hypotheical protein DKFZp564F013 NM_007173 1.46
protease, serine, 23 NM_014035 1.46
SBB31 protein AF041261 1.46
leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 4 NM_003441 1.46
zinc finger protein 141 (clone pHZ-44) NM_014962 1.46
KIAA0952 protein AA365670 1.46
Consensus includes gb:AA365670 /FEA=EST /DB_XREF=gi:2018010 /DB_XREF=est:EST76509 /UG=Hs.261526 ESTs 1.46
Consensus includes gb:AA444520 /FEA=EST /DB_XREF=gi:6986282 /DB_XREF=est:UJH-BIG-akb-h-11-O.U.I.s /CLONE=IMAGE:2733933 /UG=Hs.256986 ESTs, Moder. AA444520 1.46
vesicle trafficking, beach and anchor containing A1659561 1.46
KIAA0731 protein BC001480 1.46
leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 NM_022873 1.46
interferon, alpha-inducible protein (clone IFI-6-16) U82277 1.46
leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 2 NM_016570 1.46
CD414 AL034399 1.46
midline 2 AF017987 1.46
Consensus includes gb:AL117447.1 /DEF=Homo sapiens mRNA; cDNA DKFZp568A0617 (from clone DKFZp568A0617). /FEA=mRNA /DB_XREF=gi:5911895 /UG=Hs.30/AL117447 1.46
secreted frizzled-related protein 1 NM_002730 1.46
protein kinase, CAMP-dependent, catalytic, alpha NM_015603 1.46
DKFZP568M1019 protein NM_000029 1.46
angiotensinogen (serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 8) NM_002613 1.46
3-phosphoinositide dependent protein kinase-1 NM_0021474 1.46
Consensus includes gb:AK021474.1 /DEF=Homo sapiens cDNA FLJ11412 fs, clone HEMBA1000876. /FEA=mRNA /DB_XREF=gi:10432666 /UG=Hs.289094 Homo sapi 1.46
hypotheical protein NM_016449 1.46
recombination activating gene 1 NM_000448 1.46
regulated in glioma NM_006394 1.46
CD58 antigen, (lymphocyte function-associated antigen 3) BC005930 1.46
engulfment and cell motility 1 (ced-12 homolog, C. elegans) NM_014800 1.46
platelet activating receptor homolog NM_013308 1.46
hypotheical protein FLJ12650 NM_024522 1.46
POU domain, class 4, transcription factor 1 L20433 1.46
fracture callus 1 homolog (rat) NM_012192 1.46
cotector required for Sp1 transcriptional activation, subunit 3 (130kD) NM_004830 1.46
protein kinase, Y-linked NM_002760 1.46
aldehyde dehydrogenase 7 family, member A1 AU149534 1.46
hippocampal-like protein 4 BE550384 1.46
transient receptor potential channel 3 U47050 1.46
Cluster Inc. AF070618/Homo sapiens clone 24627 mRNA sequence /cds=UNKNOWN /gb=AF070618 /lg=3283884 /ug=Hs.14074 /len=1179 1.46
potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3 AJ251016 1.46
Consensus includes gb:AF203728.1 /DEF=Homo sapiens bone morphogenetic protein BMPY (BMPY) mRNA, partial cds. /FEA=mRNA /GEN=BMPY /PROD=bone morpho 1.46
cytochrome P450, subfamily IIIA (nhipedipine oxidase), polypeptide 3 NM_000776 1.46
lipase protein BC001698 1.46
Consensus includes gb:AK023798.1 /DEF=Homo sapiens cDNA FLJ13736 fs, clone PLACE3000156. /FEA=mRNA /DB_XREF=gi:10435842 /UG=Hs.31532 Homo sapiens 1.46
Niemann-Pick disease, type C1 NM_000271 1.46
transcription factor 7-like 2 (T-cell specific, HMG-box) A1703074 1.46
Consensus includes gb:AL050388.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564M2422 (from clone DKFZp564M2422); partial cds. /FEA=mRNA /GEN=DKFZp564M2422; 1.46
SRY (sex determining region Y)-box 4 NM_003107 1.46
sodium channel, voltage-gated, type I, alpha polypeptide AF225985 1.46
retinoic acid- and interferon-inducible protein (58kD) NM_012420 1.46
LIM domain only 1 (mombolin 1) NM_002315 1.46
folate hydrolase (prostate-specific membrane antigen) 1 NM_004476 1.46
cholinergic receptor, nicotinic, alpha polypeptide 3 K37981 1.46
ubiquitin 2 AK001029 1.46
hypotheical protein FLJ20813 NM_017961 1.46
MUF1 protein NM_006369 1.46
hypotheical protein MGC5384 NM_030972 1.46
lipase protein AA209239 1.46
ribose 5-phosphate isomerase A (ribose 5-phosphate epimerase) A1692341 1.46
KIAA0870 protein/acinus NM_014977 1.46
H3K4-B protein AF070592 1.46
pregnancy-associated plasma protein-E AF311940 1.46

Consensus includes gb:Z84717.1 /DEF=Human DNA sequence from cDNA 16pHOG:19 from chromosome 16p13.3 /FEA=mRNA /DB_XREF=gi:1817571 /UG=Hs.305284 Z84717
 xylosyltransferase II NM_022167 1.46
 empty spiracles homolog 2 (Drosophila) AF118071 1.46
 PRO1693 protein L47208 1.46
 potassium inwardly-rectifying channel, subfamily J, member 5 NM_025048 1.46
 hypothetical protein FLJ22684 NM_006200 1.46
 proprotein convertase subtilisin/kexin type 5 NM_018836 1.46
 hypothetical protein M13594 1.46
 B-cell CLL/lymphoma 2 AL021786 1.46
 integral membrane protein 2A AB014564 1.46
 KIAA0664 protein NM_014204 1.46
 BCL2-related ovarian killer NM_022564 1.46
 matrix metalloproteinase 16 (membrane-inserted) NM_008891 1.46
 crystallin, gamma D NM_031277 1.46
 ring finger protein 17 Consensus includes gb:AU147851 /FEA=EST /DB_XREF=gi:11009372 /DB_XREF=est:AU147851 /CLONE=MAMMA100187 /UG=Hs.183819 Homo sapiens cDNA FLJ12 AU147851 1.46
 integrin, beta 6 NM_000888 1.46
 hypothetical protein FLJ20241 AK023399 1.46
 spondin 1, (I-spondin) extracellular matrix protein A1885290 1.46
 zinc finger protein 146 NM_007145 1.46
 hypothetical protein FLJ10783 NM_018217 1.46
 integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41B) NM_000419 1.46
 Consensus includes gb:AV726900 /FEA=EST /DB_XREF=gi:10836321 /DB_XREF=est:AV726900 /CLONE=HTCABD05 /UG=Hs.7000 ESTs AV726900 1.46
 erythropoietin AF053356 1.46
 KIAA0644 gene product NM_014817 1.46
 protocadherin 1 (cadherin-like 1) AA461656 1.46
 BTG family, member 3 NM_008806 1.46
 v-rat simian leukemia viral oncogene homolog A (ras related) NM_005402 1.46
 hypothetical protein PRO2822 NM_018539 1.46
 inhibitor of growth family, member 1-like NM_001564 1.46
 nuclear antigen Sp100 NM_003113 1.46
 Consensus includes gb:AW968210 /FEA=EST /DB_XREF=gi:8158170 /DB_XREF=est:EST380405 /UG=Hs.293957 ESTs, Weakly similar to ALUC_HUMAN !!! ALU CLAS AW968210 1.46
 immunoglobulin lambda joining 3 AF234254 1.46
 T cell receptor delta locus X06557 1.46
 KIAA1356 protein AF352026 1.45
 immunoglobulin lambda-like polypeptide 2 L02326 1.45
 Consensus includes gb:AW970948 /FEA=EST /DB_XREF=gi:8160793 /DB_XREF=est:EST383031 /UG=Hs.269403 ESTs AW970948 1.45
 solute carrier family 18 (monocarboxylic acid transporters), member 7 AF048608 1.45
 homolog of Xenopus Claspin NM_022111 1.45
 claudin 8 AL049977 1.45
 ecdonucleotide pyrophosphatase/phosphodiesterase 2 (autolysin) D45421 1.45
 6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 4 NM_004587 1.45
 protein with polyglutamine repeat, calcium (ca2+) homeostasis endoplasmic reticulum protein NM_006387 1.45
 Consensus includes gb:A1886462 /FEA=EST /DB_XREF=gi:5591646 /DB_XREF=est:wn39e07.x1 /CLONE=IMAGE:2447844 /UG=Hs.238750 ESTs, Highly similar to serine M21191 1.45
 aldolase A, fructose-bisphosphate pseudogene 2 NM_018413 1.45
 carboxypeptidase B2 (plasma, carboxypeptidase U) Consensus includes gb:U00956.1 /DEF=Human clone KDB5.1 (CAC)(n)(GTG)n repeat-containing mRNA, /FEA=mRNA /DB_XREF=gi:405058 /UG=Hs.104112 Human clone U00956 1.45
 Consensus includes gb:AL049328.1 /DEF=Homo sapiens mRNA, cDNA DKFZp564E026 (from clone DKFZp564E026), /FEA=mRNA /DB_XREF=gi:4500102 /UG=Hs.1356 AL049328 1.45
 plasmin 1 (I isoform) NM_002670 1.45
 potassium voltage-gated channel, Shal-related subfamily, member 3 AF120491 1.45
 hypothetical protein FLJ10815 AK022786 1.45
 putative peroxisome microbody protein 175.1 NM_015887 1.45
 KIAA1361 protein AK024376 1.45
 MAD, mothers against decapentaplegic homolog 8 (Drosophila) A1628464 1.45
 PRO1398 protein NM_031289 1.45
 tight junction protein 1 (zona occludens 1) AA813018 1.45
 receptor-interacting serine-threonine kinase 2 AF027706 1.45
 cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 7 NM_000784 1.45
 T-box 6 AK022330 1.45
 ameloblastin, enamel matrix protein NM_018519 1.45
 AT2 receptor-interacting protein 1 AL098842 1.45
 CD3-epitope-associated protein; antisense to ERCC-1 NM_012099 1.45
 myristoylated alanine-rich protein kinase C substrate M88956 1.45

camitine O-octanoyltransferase
 CD84 antigen (leukocyte antigen)
 tumor necrosis factor, alpha-induced protein 6
 glutamate receptor, ionotropic, AMPA 1
 UDP glycosyltransferase 2 family, polypeptide B15
 regulatory factor X, 5 (influences HLA class II expression)
 HSPC073 protein
 hypothetical protein FLJ11292
 KIAA0892 protein
 collagen, type XI, alpha 1
 granzyme K (serine protease, granzyme 3; lyptase II)
 hypothetical protein FLJ22686
 Consensus includes gb:AJ275371 /DEF=Homo sapiens partial IGTVH3 gene for immunoglobulin heavy chain V region, case 1, clone 16 /FEA=CDS /DB_XREF=gi:7573002 /AJ275371
 GAP-associated tyrosine phosphoprotein p82 (Sam68)
 Consensus includes gb:AL050042.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566L0824 (from clone DKFZp566L0824) /FEA=mRNA /DB_XREF=gi:4894284 /UG=Hs.306 /AL050042
 ATP-binding cassette, sub-family B (MDR/TAP), member 9
 Consensus includes gb:K00827.1 /DEF=human kpn1 repeat mma (cdna clone pcd-kpn1-8), 3 end. /FEA=mRNA /DB_XREF=gi:337653 /UG=Hs.203776 Human kpn1 repeat /K00827
 solute carrier family 37 (glycerol-3-phosphate transporter), member 1
 hypothetical protein MGC20727
 LIM protein (similar to rat protein kinase C-binding enigma)
 cell recognition molecule CASPR3
 Consensus includes gb:AK025882.1 /DEF=Homo sapiens cDNA: FLJ22209 fts, clone HRC01496 /FEA=mRNA /DB_XREF=gi:10438505 /UG=Hs.288528 Homo sapiens cDNA: FLJ22209 fts, clone HRC01496 /FEA=mRNA /DB_XREF=gi:10438505 /UG=Hs.331207 ESTs
 Bcl-2 binding component 3
 KIAA0352 gene product
 paraneoplastic antigen
 hypothetical protein FLJ13942
 KIAA0766 gene product
 solute carrier family 30 (zinc transporter), member 4
 H4 histone family, member H
 hypothetical protein FLJ23282
 Consensus includes gb:BF222916 /FEA=EST /DB_XREF=gi:11130093 /DB_XREF=est:q25a04.x1 /CLONE=IMAGE:3699342 /UG=Hs.331207 ESTs
 hypothetical protein FLJ20481
 diacylglycerol kinase, beta (90kD)
 hypothetical protein FLJ11155
 azoospermia factor 1
 cAMP responsive element binding protein 1
 cyclic nucleotide-gated channel beta 1
 activating transcription factor 7
 cysteine-rich, angiogenic inducer, 61
 Consensus includes gb:AL133618.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C2021 (from clone DKFZp434C2021) /FEA=mRNA /DB_XREF=gi:6599233 /UG=Hs.30 /AL133618
 opioid receptor, mu 1
 tripartite motif-containing 29
 hypothetical protein MGC16714
 NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASH1)
 hypothetical protein FLJ21596
 adrenergic, beta-3-, receptor
 Consensus includes gb:AL050370.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566C0546 (from clone DKFZp566C0546) /FEA=mRNA /DB_XREF=gi:4914605 /UG=Hs.59 /AL050370
 Consensus includes gb:BC004857.1 /DEF=Homo sapiens, clone IMAGE:3690478, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3690478) /DB_XF BC004857
 TNF receptor-associated factor 3
 mitogen-activated protein kinase 8 interacting protein 3
 potassium voltage-gated channel, shaker-related subfamily, member 10
 homeo box A9
 Consensus includes gb:AL049798 /DEF=Human DNA sequence from clone 797M17 on chromosome 1q22-24.3. Contains the DPT gene for Dermatopontin, ESTs, en STS /AL049798
 PR domain containing 2, with ZNF domain
 class I cytoline receptor
 potassium voltage-gated channel, Shaw-related subfamily, member 4
 calpain 3, acidic
 aquaporin 4
 dead ringer-like 1 (Drosophila)
 RAN binding protein 3
 gb:NM_012348.1 /DEF=Homo sapiens olfactory receptor 89 (OLFMR89), mRNA. /FEA=CDS /GEN=OLFMR89 /PROD=olfactory receptor 89 /DB_XREF=gi:6912547 /UG=Hs.2 /NM_012348
 Sjogren syndrome antigen A2 (SjokD, ribonucleoprotein autoantigen SS-A/Ro)

NM_021151 1.45
 AF054816 1.45
 NM_007115 1.45
 M64752 1.45
 U08841 1.45
 AW027312 1.45
 NM_014163 1.45
 AK023539 1.45
 AC003030 1.45
 NM_001854 1.45
 NM_002104 1.45
 AB25877 1.45
 AW592227 1.45
 AW592227 1.45
 AK01040 1.45
 BE504895 1.45
 NM_018984 1.45
 AF131745 1.45
 BF671400 1.45
 NM_024879 1.45
 AK025882 1.45
 AF332558 1.45
 NM_014830 1.45
 AL22335 1.45
 NM_024581 1.45
 NM_014805 1.45
 NM_013309 1.45
 NM_003543 1.45
 NM_024816 1.45
 BF222916 1.45
 NM_017833 1.45
 AB018261 1.45
 NM_018342 1.45
 AC007320 1.45
 NM_004379 1.45
 U16945 1.45
 NM_006856 1.45
 NM_001554 1.45
 AL133618 1.45
 NM_000914 1.45
 AF230389 1.45
 AL117502 1.45
 NM_024823 1.45
 AA723057 1.45
 NM_000025 1.45
 AL050370 1.45
 BC004857 1.45
 NM_003300 1.45
 AB028689 1.45
 NM_005549 1.45
 AL246769 1.45
 AL049798 1.45
 U23736 1.45
 AB931115 1.45
 NM_004978 1.45
 NM_001839 1.45
 D83412 1.45
 NM_005224 1.45
 BC004349 1.45
 NM_012348 1.45
 AK024044 1.45

testicular hspidol expressed gene	NM_016585	1.45
phospholipase A2, group IID	AL158172	1.45
GATA binding protein 1 (globin transcription factor 1)	M30601	1.45
Ras homolog enriched in brain 2	D78132	1.45
hypothetical protein MGC13138	AA746290	1.45
ribosomal protein L18	AI665349	1.45
gb:NM_003723.1 /DEF=Homo sapiens caspase 13, apoptosis-related cysteine protease (CASP13), mRNA. /FEA=mRNA /GEN=CASP13 /PROD=caspase 13 /DB_XREF=	NM_003723	1.45
ribosomal protein S6 kinase, 90kD, polypeptide 6	NM_014496	1.45
mitogen-activated protein kinase kinase 1 interacting protein 1	NM_021970	1.45
similar to slyltransferase 7 (alpha-N-acetylneuraminyl-2,3-betagalactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase) E	NM_030965	1.45
pregnancy-induced growth inhibitor	NM_013370	1.45
hypothetical protein FLJ21901	NM_024622	1.45
tumor necrosis factor (ligand) superfamily, member 13	AF114011	1.45
hypothetical protein FLJ23263	NM_025115	1.45
Apobec-1 complementation factor; APOBEC-1 stimulating protein	NM_014576	1.45
CMRF35 leukocyte immunoglobulin-like receptor	NM_006678	1.45
serine (or cysteine) proteinase inhibitor, clade F (alpha-2 antipain, pigment epithelium derived factor), member 2	NM_000934	1.45
leukemia inhibitory factor receptor	NM_002310	1.45
Consensus includes gb:BF433429 /FEA=EST /DB_XREF=gi:11445592 /DB_XREF=est:7q53a12.x1 /CLONE=IMAGE:3702046 /UG=Hs.289068 Homo sapiens cDNA FLJ11 BF433429	1.45	
Consensus includes gb:U79265.1 /DEF=Human clone 23614 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710224 /UG=Hs.91375 Human clone 23614 mRNA sequence U79265	1.45	
Consensus includes gb:U05004.1 /DEF=Homo sapiens mRNA; cDNA DKFpZp568M0524 (from clone DKFpZp568M0524). /FEA=mRNA /DB_XREF=gi:4884285 /UG=Hs.3C AL050043	1.45	
Consensus includes gb:D29810.1 /DEF=Human mRNA for unknown product, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:704440 /UG=Hs.153445 Human mf D29810	1.45	
Rev/Rex activation domain binding protein-related	AF015042	1.45
Consensus includes gb:M16917 /DEF=Human glucocerebrosidase gene 10-2, 5 and /FEA=CDS_1 /DB_XREF=gi:183025 /UG=Hs.287396 Human glucocerebrosidase gene M16917	1.45	
ZYG homolog	BE675849	1.45
nephronophthisis 1 (juvenile)	NM_000272	1.45
lymphotxin alpha (TNF superfamily, member 1)	NM_000595	1.45
Consensus includes gb:A017119 /FEA=EST /DB_XREF=gi:3231455 /DB_XREF=est:ov34a05.x1 /CLONE=IMAGE:1639184 /UG=Hs.144339 Human DNA sequence from c A017119	1.45	
Consensus includes gb:AU143984 /FEA=EST /DB_XREF=gi:11005505 /DB_XREF=est:AU143984 /CLONE=HEMBA1000568 /UG=Hs.301645 Homo sapiens cDNA FLJ100 AU143984	1.45	
glutamate receptor, metabotropic 3	NM_000840	1.45
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson AB030075	1.45	
Cluster incl. A937160/vp3f02.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2467443 /clone_end=3' /gb=A937160 /gi=5676030 /ug=Hs.7967 /len=552	1.45	
senum constituent protein	NM_007061	1.45
nucleic diphosphate linked moiety X-type motif 7	AK026469	1.45
glycophorin A (includes MN blood group)	U00178	1.45
Consensus includes gb:AK024185.1 /DEF=Homo sapiens cDNA FLJ14123 fts, clone MAMMA1002155. /FEA=mRNA /DB_XREF=gi:10436502 /UG=Hs.269314 Homo sapiens AK024185	1.45	
carbamoyl-phosphate synthetase 1, mitochondrial	AF154830	1.45
slatherin	NM_003154	1.45
gb:U05102.1 /DEF=Human IG processed VH-region mRNA, complete cds. /FEA=mRNA /GEN=IGH@ /PROD=immunoglobulin heavy chain V region /DB_XREF=gi:185528 /L06102	1.45	
Consensus includes gb:AK025143.1 /DEF=Homo sapiens cDNA: FLJ21490 fts, clone COL05464. /FEA=mRNA /DB_XREF=gi:10437602 /UG=Hs.288700 Homo sapiens cC AK025143	1.44	
postmeiotic segregation increased 2-like 6	D39503	1.44
steroidogenic acute regulatory protein	NM_000349	1.44
fer-1-like 3, myofibrin (C. elegans)	NM_013451	1.44
prostaglandin E synthase	NM_004878	1.44
Consensus includes gb:AL079289.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 35971. /FEA=mRNA /DB_XREF=gi:5102748 /UG=Hs.137154 /AL079289	1.44	
hypothetical protein FLJ10675	NM_018176	1.44
intersectin 1 (SH3 domain protein)	AF114488	1.44
serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1	NM_000602	1.44
KIAA1037 protein	AB028960	1.44
lyyyl oxidase	BE503425	1.44
glutamate receptor, ionotropic, N-methyl D-aspartate 1	AF015731	1.44
lms-related tyrosine kinase 1 (vascular endothelial growth factor/vascular permeability factor receptor)	AA058828	1.44
killer cell lectin-like receptor subfamily D, member 1	U30610	1.44
Consensus includes gb:AF052140.1 /DEF=Homo sapiens clone 23758 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360449 /UG=Hs.141055 Homo sapiens clone 23758 AF052140	1.44	
Consensus includes gb:X63118 /DEF=H. sapiens DNA for oestrogen receptor gene promoter /FEA=CDS /DB_XREF=gi:351159 /UG=Hs.247938 H. sapiens DNA for oestrogen X63118	1.44	
Mahlavu hepatocellular carcinoma	NM_006543	1.44
transcription factor AP-2 beta (activating enhancer binding protein 2 beta)	NM_003221	1.44
transmembrane 4 superfamily member (tetraspan NET-2)	NM_012338	1.44
hypothetical protein FLJ20950	NM_024952	1.44
Consensus includes gb:BF110551 /FEA=EST /DB_XREF=gi:10940241 /DB_XREF=est:7n38h09.x1 /CLONE=IMAGE:3567208 /UG=Hs.253600 ESTs, Highly similar to S50 BF110551	1.44	
insulin-like growth factor binding protein 5	AW157548	1.44
lectin med-2	NM_014921	1.44
solute carrier family 14 (urea transporter), member 1 (Kidd blood group)	NM_015865	1.44

KIAA1030 protein	AB028953	1.44
EphA5	L36844	1.44
SA (rat hypertension-associated) homolog	D16350	1.44
ferrochelatase (protoporphyrin)	NM_000140	1.44
kinesin family member C3	BC001211	1.44
JM1 protein	BF224247	1.44
netrin 2-like (chicken)	NM_006181	1.44
hepatocellular carcinoma novel gene-3 protein	NM_016651	1.44
hypothetical protein FLJ23534	NM_024806	1.44
solute carrier family 4, sodium bicarbonate cotransporter, member 4	AF157492	1.44
ets variant gene 6 (TEL oncogene)	AF041811	1.44
hypothetical protein PRO1895	NM_016812	1.44
hypothetical protein FLJ20337	AL096712	1.44
splicing factor, arginine/serine-rich 11	AU146237	1.44
phosphatidylinositol glycan, class K	AF022813	1.44
hypothetical protein FLJ10786	NM_018219	1.44
B-cell CLL/lymphoma 11A (zinc finger protein)	NM_022893	1.44
empty spiracles homolog 1 (Drosophila)	BF515060	1.44
mucin 5, subtype B, tracheobronchial	AI521646	1.44
holocytochrome c synthase (cytochrome c heme-lyase)	AI801013	1.44
Consensus includes gb:AW974995 /FEA=EST /DB_XREF=estEST387100 /UG=Hs.292787 ESTs	AW974995	1.44
KIAA0699 protein	AI934125	1.44
KIAA1822 protein	BE300056	1.44
killer cell lectin-like receptor subfamily D, member 1	AB009597	1.44
forkhead box E1 (thyroid transcription factor 2)	NM_004473	1.44
hypothetical protein FLJ22031	NM_025074	1.44
Consensus includes gb:AF090910.1 /DEF=Homo sapiens clone HQ0262 /FEA=mRNA /DB_XREF=gi:5690195 /UG=Hs.236760 Homo sapiens clone HQ0262	AF090910	1.44
Consensus includes gb:AK026439.1 /DEF=Homo sapiens cDNA: FLJ22786 fls, clone KAA2150, /FEA=mRNA /DB_XREF=gi:10439304 /UG=Hs.306848 Homo sapiens cDI	AK022046	1.44
kallikrein 14	NM_000037	1.44
ankyrin 1, erythrocytic	AW615812	1.44
ATPase, Ca++ transporting, plasma membrane 3	NM_005369	1.44
MCF.2 cell line derived transforming sequence	H19843	1.44
synapsin I	NM_017937	1.44
hypothetical protein FLJ20712	AU150824	1.44
DiGeorge syndrome critical region gene 2	NM_003579	1.44
RAD54-like (S.cerevisiae)	NM_006604	1.44
ret finger protein-like 3	NM_005019	1.44
phosphodiesterase 1A, calmodulin-dependent	AB004903	1.44
STAT induced STAT inhibitor-2	D13515	1.44
glutamate receptor, ionotropic, N-methyl D-aspartate 1	NM_024886	1.44
cytochrome P450, subfamily IVF, polypeptide 8	AF133296	1.44
adaptor-related protein complex 3, sigma 1 subunit	NM_001284	1.44
proprotein convertase subtilisin/kexin type 2	NM_002594	1.44
nuclear receptor subfamily 5, group A, member 1	U76388	1.44
neuronal thread protein	AF010144	1.44
solute carrier family 21 (organic anion transporter), member 3	NM_021094	1.44
disabled homolog 1 (Drosophila)	NM_021080	1.44
microtubule-associated protein 1A	AW298788	1.44
matrix metalloproteinase 17 (membrane-inserted)	NM_016155	1.44
homeo box A3	NM_030661	1.44
heterogeneous nuclear ribonucleoprotein H1 (H)	AI022387	1.44
glycogen synthase 2 (liver)	S70004	1.44
NGF-A binding protein 1 (EGR1 binding protein 1)	AF045451	1.44
Fc fragment of IgG, high affinity Ia, receptor for (CD64)	X14355	1.44
UDP-Gal:betaGlcNAc beta 1.4-galactosyltransferase, polypeptide 2	BC002431	1.44
phosphodiesterase 3A, cGMP-inhibited	U36788	1.44
hypothetical protein MGC4643	AA824389	1.44
anaplastic lymphoma kinase (K1-1)	BF057492	1.44
adenylyate cyclase 8	AB011092	1.44
aggrekan 1 (chondrolectin sulfate proteoglycan, antigen identified by monoclonal antibody A0122)	AF349314	1.44
Zn family member 1 (odd-paired homolog, Drosophila)	NM_001135	1.44
	NM_003412	1.44

fms-related tyrosine kinase 1 (vascular endothelial growth factor/vascular permeability factor receptor)
 MCN3 minichromosome maintenance deficient 3 (S. cerevisiae) associated protein
 homolog of yeast ATP12
 Consensus includes gb:AK022038.1 /DEF=Homo sapiens cDNA FLJ11976 fls, clone HEMBB1001253. /FEA=mRNA /DB_XREF=gi:10433356 /UG=Hs.296879 Homo sapie
 KIAA0063 gene product
 hypothetical protein R32184_1
 solute carrier family 5 (neutral amino acid transporters, system A), member 4
 phosphoinositide-3-kinase, catalytic, beta polypeptide
 eve, even-skipped homeo box homolog 1 (Drosophila)
 RAD51-like 1 (S. cerevisiae)
 KIAA0318 protein
 Consensus includes gb:S80491.1 /DEF=stem cell factor (alternatively spliced) human, preimplantation embryos, blastocysts, mRNA Partial, 180 nt. /FEA=mRNA /GEN=ste
 cadherin 15, M-cadherin (myotubule)
 T-box 1
 Consensus includes gb:AK024393.1 /DEF=Homo sapiens cDNA FLJ14331 fls, clone PLACE4000320. /FEA=mRNA /DB_XREF=gi:10436771 /UG=Hs.288569 Homo sapie
 KIAA0977 protein
 tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain
 solute carrier family 18 (folate transporter), member 1
 mitochondrial ribosomal protein L33
 Consensus includes gb:AL035295.1 /DEF=H. sapiens gene from PAC 106H8, similar to Dynamin. /FEA=mRNA /PROD=hypothetical protein /DB_XREF=gi:4200217 /UG=Hs
 hypothetical protein DKFZ0761K1824
 likely homolog of mouse heparin/heparin sulfate-glucuronidase C5-epimerase
 hypothetical protein DKFZp761K1423
 adenylyl cyclase 2 (brain)
 Consensus includes gb:AK024355.1 /DEF=Homo sapiens cDNA FLJ14293 fls, clone PLACE1007866. /FEA=mRNA /DB_XREF=gi:10436719 /UG=Hs.141009 Homo sapie
 hypothetical protein OVC10-2
 hypothetical protein FLJ13941
 KIAA0672 gene product
 myeloid cell nuclear differentiation antigen
 melanoma cell adhesion molecule
 Lin-7b protein; likely ortholog of mouse LIN-7B; mammalian LIN-7 protein 2
 nuclear fragile X mental retardation protein interacting protein 1
 HSPC048 protein
 Consensus includes gb:AK023187.1 /DEF=Homo sapiens cDNA FLJ13125 fls, clone NT2RP3002877. /FEA=mRNA /DB_XREF=gi:10435001 /UG=Hs.287551 Homo sapie
 chloride channel 1, skeletal muscle (Thomsen disease, autosomal dominant)
 TIA1 cytotoxic granule-associated RNA binding protein
 zinc finger protein 108
 calmodulin 2 (phosphorylase kinase, delta)
 mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme B
 KIAA0485 protein
 Consensus includes gb:AF052173.1 /DEF=Homo sapiens clone 24417 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360484 /UG=Hs.306253 Homo sapiens clone 24417 AF052173
 Consensus includes gb:BF002254. /FEA=EST /DB_XREF=gi:10702529 /DB_XREF=est:7h02a03.x1 /CLONE=IMAGE.3314764 /UG=Hs.143600 type II Golgi membrane pro
 KIAA0680 gene product
 midline 1 (Optiz/BBB syndrome)
 B-cell CLL/lymphoma 10
 integrin, beta 3 (platelet glycoprotein IIb, antigen CD61)
 discs, large (Drosophila) homolog 2 (chapsyn-110)
 hypothetical protein FLJ10043
 Consensus includes gb:N64802 /FEA=EST /DB_XREF=gi:1212631 /DB_XREF=est:y231b05.s1 /CLONE=IMAGE.284625 /UG=Hs.173684 Homo sapiens mRNA; cDNA DK1 N64802
 Consensus includes gb:AI206888 /FEA=EST /DB_XREF=gi:3765560 /DB_XREF=est:q35a04.x1 /CLONE=IMAGE.1942830 /UG=Hs.154131 ESTs
 tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)
 hypothetical protein FLJ22477
 D15F37 (pseudogene)
 Consensus includes gb:AL049442.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586N1720 (from clone DKFZp586N1720). /FEA=mRNA /DB_XREF=gi:4500222 /UG=Hs.71
 phosphatidylinositol-4-phosphate 5-kinase, type I, beta
 myeloid/lymphoid or mixed-lineage leukemia (rithorax homolog, Drosophila); translocated to, 2
 hypothetical protein FLJ20105
 four and a half LIM domains 1
 hypothetical protein FLJ11113
 disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)
 host cell factor C1 (VP16-accessory protein)
 MD-2 protein
 lethal giant larvae homolog 1 (Drosophila)

allograft inflammatory factor 1
 olfactory receptor, family 1, subfamily K, member 1
 arylsulfatase A
 hematopoietic PBX-interacting protein
 glycoprotein M6B
 hypothetical protein PRO1728
 Consensus includes gb:L37198.1 /DEF=Homo sapiens (clone B3B3E13) Huntingtons disease candidate region mRNA fragment. /FEA=mRNA /DB_XREF=gi:600519 /UG=L37198
 dickkopf homolog 3 (Xenopus laevis)
 Consensus includes gb:AK026667.1 /DEF=Homo sapiens cDNA: FLJ23014 fis, clone LNG00806. /FEA=mRNA /DB_XREF=gi:10439568 /UG=Hs.306862 Homo sapiens cDNA: AK026667
 hypothetical protein FLJ10761
 Consensus includes gb:AK022442.1 /DEF=Homo sapiens cDNA: FLJ12380 is, clone MAMMA1002556. /FEA=mRNA /DB_XREF=gi:10433838 /UG=Hs.300934 Homo sapiens cDNA: AK022442
 v-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian)
 forhead box L1
 synaptophysin, gamma 2
 DLEU2L
 chromosome 4 open reading frame 5
 Consensus includes gb:AL110227.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434J194 (from clone DKFZp434J194). /FEA=mRNA /DB_XREF=gi:5817165 /UG=Hs.32306 AL110227
 KIAA1305 protein
 Consensus includes gb:AL030995 /DEF=Human DNA sequence from clone 1170D6 on chromosome Xq22.3-23. Contains a pseudogene similar to U-SNRNP-associated (AL030995)
 Consensus includes gb:AL030995 /DEF=Human DNA sequence from clone 1170D6 on chromosome Xq22.3-23. Contains a pseudogene similar to U-SNRNP-associated (AL030995)
 Notch homolog 2 (Drosophila)
 KIAA0257 protein
 Consensus includes gb:AL157902 /DEF=Human DNA sequence from clone RPA-675C20 on chromosome 1p13.2. Contains the 3' end of the MAN1A2 gene for mannosidase AL157902
 major histocompatibility complex, class II, DP alpha 1
 X17013 B subtilis lys gene for diaminopimelate decarboxylase corresponding to nucleotides 350-1345 of X17013 (-5, -M, -3 represent transcript regions 5 prime, Middle, ar X17013
 betacellulin
 Consensus includes gb:AL133396 /DEF=Human DNA sequence from clone RPS-1068H6 on chromosome 20q11.1-11.23. Contains a pseudogene similar to D11 (isopentate) AL133396
 Consensus includes gb:AW675655 /FEA=EST /DB_XREF=gi:7540890 /DB_XREF=est:ba52601.x1 /CLONE=IMAGE:2900184 /UG=Hs.314158 ESTs
 nucleoporin 62KD
 TAF₉-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65 kD
 undifferentiated embryonic cell transcription factor 1
 Consensus includes gb:AL035555 /DEF=Human DNA sequence from clone 217P22 on chromosome 6p21.1-21.31. Contains part of a novel Dynein heavy chain family prot AL035555
 retinoblastoma binding protein 2
 ATP-binding cassette, sub-family A (ABC1), member 3
 Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse)
 calcium channel, voltage-dependent, alpha 2/delta subunit 2
 radixin
 calcium/calmodulin-dependent protein kinase (CaM kinase) II alpha
 hypothetical protein, estradiol-induced
 galactosidase, alpha
 peroxisomal biogenesis factor 3
 serine racemase
 lipin 1
 hypothetical protein FLJ20069
 Consensus includes gb:AL133018.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434F0327 (from clone DKFZp434F0327). /FEA=mRNA /DB_XREF=gi:6453498 /UG=Hs.271 AL133018
 small optic tectal homolog (Drosophila)
 Consensus includes gb:BE658194 /FEA=EST /DB_XREF=gi:10372818 /DB_XREF=est:7g19c05.x1 /CLONE=IMAGE:3306920 /UG=Hs.323053 Homo sapiens mRNA full le BE658194
 potassium inwardly-rectifying channel, subfamily J, member 14
 phosphodiesterase 7B
 hyaluronoglucosaminidase 1
 brain-specific angiogenesis inhibitor 3
 HT001 protein
 ankyrin repeat domain 2 (stretch responsive muscle)
 calcium binding protein 1 (calbrain)
 forhead box C2 (MPH-1, mesenchyme forhead 1)
 hypothetical protein DKFZp781B2423
 heat shock transcription factor 2 binding protein
 Consensus includes gb:A478300 /FEA=EST /DB_XREF=gi:4371526 /DB_XREF=est:tm39601.x1 /CLONE=IMAGE:2160504 /UG=Hs.192789 ESTs, Weakly similar to ALUK A478300
 Dlx-binding protein IDAX (inhibition of the Dlx and Axin complex)
 ATPase, Ca++ transporting, type 2C, member 1
 deleted in lymphocytic leukemia, 2
 solute carrier family 6 (neurotransmitter transporter, glycine), member 5
 phosphodiesterase 4D interacting protein (myomegalin)

wingless-type MMTV integration site family, member 108
hypothetical protein FLJ22595
Consensus includes gb:AL110248.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434N021 (from clone DKFZp434N021) /FEA=mRNA /DB_XREF=gi:5817198 /UG=Hs.2087
protocadherin beta 11
epsin 3
haptoglobin-related protein
alpha-catenin-like protein
cell cycle progression 8 protein
lactylkinin, precursor 1 (substance K, neurokinin 1, neurokinin 2, neuromedin L, neurokinin alpha, neuropeptide K, neuropeptide gamma)
paired box gene 3 (Wardenburg syndrome 1)
hypothetical protein FLJ14639
arginine vasopressin receptor 2 (nephrogenic diabetes insipidus)
fetuin B
ras-like protein
hypothetical protein
hypothetical protein FLJ13576
tumor necrosis factor (ligand) superfamily, member 12
Consensus includes gb:AL049430.1 /DEF=Homo sapiens mRNA; cDNA DKFZp568H201 (from clone DKFZp568H201) /FEA=mRNA /DB_XREF=gi:4500210 /UG=Hs.3020
ectodermal dysplasia 1, anhidrotic
ring finger protein 4
protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calciuretin A gamma)
serine protease inhibitor, Kunitz type, 3
hypothetical protein FLJ11100
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 4
S164 protein
Consensus includes gb:AK024445.1 /DEF=Homo sapiens mRNA for FLJ00035 protein, partial cds. /FEA=mRNA /GEN=FLJ00035 /PROD=FLJ00035 protein /DB_XREF=gi:AK024445
stailidase 3 (membrane sialidase)
protein kinase, cGMP-dependent, type II
opioid receptor, mu 1
E2F transcription factor 5, p130-binding
solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2
Consensus includes gb:AW450360 /FEA=EST /DB_XREF=gi:6991136 /DB_XREF=est:UL-H-B13-akn-c-11-0-UJ.s1 /CLONE=IMAGE:2734845 /UG=Hs.98834 ESTs
KIAA0161 gene product
translocase of inner mitochondrial membrane 17 homolog A (yeast)
t-complex 1
Consensus includes gb:BE80245 /FEA=EST /DB_XREF=gi:10329021 /DB_XREF=est:601490812F1 /CLONE=IMAGE:3893018 /UG=Hs.164038 Homo sapiens AKAP350
transcription elongation factor B (SIII), polypeptide 2 (18kD, elongin B)
dual specificity phosphatase 4
GTP-binding protein Rho7
kinesin family member 13A
lengsin
pregnancy specific beta-1-glycoprotein 3
mitochondrial ribosomal protein 63
Consensus includes gb:AK024128.1 /DEF=Homo sapiens cDNA FLJ14066 fs, clone HEMBB1001197 /FEA=mRNA /DB_XREF=gi:10436433 /UG=Hs.306865 Homo sapie
neuregulin 1
LIV-1 protein, estrogen regulated
cadherin-like protein VR20
secretogranin III
neurogenin 1
nucleobindin 1
ets variant gene 1
hypothetical protein FLJ12270
glutamate receptor, ionotropic, kainate 5
hypothetical protein
GTP binding protein 1
Usher syndrome 2A (autosomal recessive, mild)
gap junction protein, alpha 8, 50kD (connexin 50)
golgln-67
Consensus includes gb:AL049331.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564I066 (from clone DKFZp564I066) /FEA=mRNA /DB_XREF=gi:4500106 /UG=Hs.207805
hypothetical protein FLJ20069
S100 calcium binding protein A6 (calicyclin)
kinesin-like 3

NM_003394 1.42
NM_025047 1.42
AL110248 1.42
NM_018931 1.42
NM_017957 1.42
NM_020995 1.42
NM_013266 1.42
NM_004748 1.42
NM_003182 1.42
U02309 1.42
AA152202 1.42
NM_000054 1.42
AB017551 1.42
BF670447 1.42
BC002587 1.42
NM_022484 1.42
NM_003809 1.42
AL049430 1.42
AF081192 1.42
BF986633 1.42
BC004864 1.42
AW118166 1.42
NM_018321 1.42
U82079 1.42
AB25305 1.42
AK024445 1.42
AK022450 1.42
NM_006259 1.42
U12569 1.42
AW139195 1.42
NM_000617 1.42
AW450360 1.42
NM_014746 1.42
BC005832 1.42
NM_006335 1.42
BF224073 1.42
BE880245 1.42
AC004493 1.42
NM_001394 1.42
BF511718 1.42
NM_022113 1.42
NM_016571 1.42
R32065 1.42
BF195165 1.42
AK024128 1.42
NM_004495 1.42
AB354449 1.42
AL109928 1.42
NM_013243 1.42
NM_008161 1.42
BC002356 1.42
X87175 1.42
NM_030581 1.42
AL137705 1.42
AA160474 1.42
NM_004286 1.42
NM_007123 1.42
NM_005267 1.42
AW008438 1.42
AL049331 1.42
NM_017851 1.42
NM_014624 1.42
NM_030615 1.42

acyltransferase (neutrophil)
 Consensus includes gb:BE742268 /FEA=EST /DB_XREF=gi:10156260 /DB_XREF=est:60157568F.1 /CLONE=IMAGE:3836709 /UG=Hs.281706 sortilin 1
 Consensus includes gb:R43279 /FEA=EST /DB_XREF=gi:821386 /DB_XREF=est:yg17c12.31 /CLONE=IMAGE:32731 /UG=Hs.22574 ESTs
 hypothetical protein FLJ11800
 Consensus includes gb:AA156865 /FEA=EST /DB_XREF=gi:1728480 /DB_XREF=est:z20008.s1 /CLONE=IMAGE:502503 /UG=Hs.22370 Homo sapiens mRNA; cDNA DK A032987
 DKFZP434N043 protein
 cyclin T2
 p8 protein (candidate of metastasis 1)
 bone morphogenetic protein 2
 solute carrier family 9 (sodium/hydrogen exchanger), isoform 3
 cyclochrome b-245, beta polypeptide (chronic granulomatous disease)
 testis intracellular mediator protein
 tollid-like 2
 Consensus includes gb:AL139318 /DEF=Human DNA sequence from clone RP11-124B17 on chromosome 13. Contains the DCT gene for dopachrome tautomerase (dopa AL139318
 Consensus includes gb:AK024945.1 /DEF=Homo sapiens cDNA; FLJ21292.fts, clone COL01869 /FEA=mRNA /DB_XREF=gi:10437367 /UG=Hs.306732 Homo sapiens cDNA AK024945
 adenylate cyclase 3
 UDP-N-acetyl-alpha-D-galactosamine-(N-acetylneuraminyl)-galactosylglucosylceramide N-acetylglucosaminyltransferase (GalNAc-T)
 transmembrane 7 superfamily member 1 (upregulated in kidney)
 guanine nucleotide exchange factor for Rap1; M-Ras-regulated GEF
 PRO0038 protein
 guanine nucleotide exchange factor for Rap1; M-Ras-regulated GEF
 protein tyrosine phosphatase, receptor type, M
 cyclin-dependent kinase inhibitor 2A (melanoma, p16, inhibits CDK4)
 phospholipase A2, group V
 nucleoporin 153KO
 parathyroid hormone receptor 1
 chitinase, di-N-acetyl-
 gliat cells missing homolog b (Drosophila)
 RAP1, GTPase activating protein 1
 PRO0457 protein
 malic enzyme 1, NAD(P)+-dependent, cytosolic
 Consensus includes gb:U06813.1 /DEF=Homo sapiens serotonin-7 receptor pseudogene, complete sequence. /FEA=mRNA /PROD=serotonin-7 receptor /DB_XREF=gi:31 U06813
 Consensus includes gb:AL571723 /FEA=EST /DB_XREF=gi:12929303 /DB_XREF=est:AL571723 /CLONE=CSODI031YD10 (3 prime) /UG=Hs.7247 ESTs, Weakly similar 1
 vesicle-associated membrane protein 3 (cellubrevin)
 tropomyosin 2 (beta)
 G protein-coupled receptor kinase 2-like (Drosophila)
 chemokine (C-C motif) receptor 3
 hypothetical protein MQC2718
 hypothetical protein FLJ20071
 1,2-alpha-mannosidase IC
 homolog of mouse quaking QKI (KH domain RNA binding protein)
 chromosome 1 open reading frame 24
 ichthy homolog E3 ubiquitin protein ligase (mouse)
 Consensus includes gb:AK024860.1 /DEF=Homo sapiens cDNA; FLJ21207.fts, clone COL00362. /FEA=mRNA /DB_XREF=gi:10437268 /UG=Hs.306708 Homo sapiens cDNA AK024860
 KIAA0443 gene product
 KIAA0831 protein
 hypothetical protein FLJ23510
 heat shock 70kD protein 1A
 interleukin 16 (lymphocyte chemoattractant factor)
 Consensus includes gb:AF055011.1 /DEF=Homo sapiens clone 24587 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3005734 /UG=Hs.137173 Homo sapiens clone 24587 AF055011
 leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 1
 FOS-like antigen 2
 collagen, type XI, alpha 1
 transcription elongation factor A (SII)-like 1
 Consensus includes gb:AL049437.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566E1120 (from clone DKFZp566E1120) /FEA=mRNA /DB_XREF=gi:4500220 /UG=Hs.10 /AL049437
 testis expressed sequence 11
 adaptor-related protein complex 1, mu 2 subunit
 phospholipase A2, group VI (cytosolic, calcium-independent)
 lymphocyte antigen 84 homolog, activating NK-receptor, NK-p48, (mouse)
 Consensus includes gb:AL110180.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J2116 (from clone DKFZp564J2116) /FEA=mRNA /DB_XREF=gi:5817104 /UG=Hs.306 /AL110180
 ras homolog gene family, member 1
 hypothetical protein FL10803
 WD repeat domain 9

laminin, alpha 3 (nican (150KD), kalinin (165KD), BM600 (150KD), epieglin)
 BTG family, member 3
 fatty acid binding protein 7, brain
 hypothetical protein FLJ20401
 deduced protein induced by progesterone
 myristoylated alanine-rich protein kinase C substrate
 Consensus includes gb:AL133075.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M0317.1 (from done DKFZp434M0317). /FEA=mRNA /DB_XREF=gi:6453518 /UG=Hs.27 AF133075
 gb:AF130051.1 /DEF=Homo sapiens clone FLB3535 PRO0898 mRNA, complete cds. /FEA=mRNA /PROD=PRO0898 /DB_XREF=gi:11493408 /UG=Hs.306860 Homo sap AF130051
 cyclokeratin 20
 hypothetical protein CG003
 putative GR6 protein
 coagulation factor II (thrombin) receptor-like 1
 CD68 antigen
 sine oculis homeobox homolog 6 (Drosophila
 allograft inflammatory factor 1
 occludin
 hypothetical protein
 hypothetical protein FLJ10833
 interleukin 1 receptor accessory protein-like 2
 deleted in liver cancer 1
 sarcoglycan, gamma (35KD dystrophin-associated glycoprotein)
 olfactory receptor, family 10, subfamily H, member 3
 Fc fragment of IgC, low affinity II, receptor for (CD23A)
 Consensus includes gb:AF009267.1 /DEF=Homo sapiens clone FBA1 Cr-du-chat region mRNA. /FEA=mRNA /DB_XREF=gi:2331069 /UG=Hs.102238 Homo sapiens dbn. AF009267
 vitelliform macular dystrophy (Best disease, bestrophin)
 cystatin A (stefin A)
 phosphatidylinositol glycan, class O
 Consensus includes gb:AW051172 /FEA=EST /DB_XREF=gi:6040324 /DB_XREF=est:tx60609.x1 /CLONE=IMAGE:2606368 /UG=Hs.243366 ESTs, Highly similar to KPCI
 serine/arginine repetitive matrix 1
 Consensus includes gb:AU148024 /FEA=EST /DB_XREF=gi:11009545 /DB_XREF=est:AU148024 /CLONE=MAMMA1002392 /UG=Hs.302075 Homo sapiens cDNA FLJ12 AU148024
 forthead box O1A (maodomyosarcoma)
 Consensus includes gb:AK021930.1 /DEF=Homo sapiens cDNA FLJ11868 fts, clone HEMBA1006993. /FEA=mRNA /DB_XREF=gi:104333231 /UG=Hs.302280 Homo saple AK021930
 KIAA1032 protein
 Consensus includes gb:L13283.1 /DEF=Homo sapiens (clone MG2-5-12) mucin (MG2) mRNA, complete polyA site. /FEA=mRNA /DB_XREF=gi:292518 /UG=Hs.103944 H.L13283
 FAT tumor suppressor homolog 1 (Drosophila)
 Consensus includes gb:AU346341 /FEA=EST /DB_XREF=gi:4083547 /DB_XREF=est:tp50b09.x1 /CLONE=IMAGE:1926425 /UG=Hs.298998 Homo sapiens, Similar to hyp AU346341
 CD36 antigen (collagen type I receptor, thrombospondin receptor)
 RCE1 homolog, prenyl protein protease (S. cerevisiae)
 retinoid acid receptor-beta associated open reading frame
 Consensus includes gb:AK024568.1 /DEF=Homo sapiens cDNA: FLJ20915 fts, clone ADSE00692. /FEA=mRNA /DB_XREF=gi:10436878 /UG=Hs.306691 Homo sapiens c AK024568
 hypothetical protein FLJ11292
 neuritin
 eukaryotic translation initiation factor 3, subunit 5 (epsilon, 47KD)
 Consensus includes gb:AT41876 /FEA=EST /DB_XREF=gi:5110251 /DB_XREF=est:wp48b05.x1 /CLONE=IMAGE:2368305 /UG=Hs.194051 Homo sapiens mRNA; cDNA AT41876
 gb:AF118715.1 /DEF=Homo sapiens PRO2829 mRNA, complete cds. /FEA=mRNA /PROD=PRO2829 /DB_XREF=gi:7959928 /UG=Hs.236256 Homo sapiens PRO2829 m AF118715
 Consensus includes gb:AK026847.1 /DEF=Homo sapiens cDNA: FLJ23194 fts, clone REC00490. /FEA=mRNA /DB_XREF=gi:10439802 /UG=Hs.306887 Homo sapiens c AK026847
 KIAA0830 protein
 guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 1
 Consensus includes gb:AW771015 /FEA=EST /DB_XREF=gi:7703066 /DB_XREF=est:hms5q005.x1 /CLONE=IMAGE:3027081 /UG=Hs.279020 Homo sapiens mRNA; cDN/ AW771015
 gb:NM_022125.1 /DEF=Homo sapiens immortalization-upregulated protein (IMUP), mRNA. /FEA=mRNA /GEN=IMUP /PROD=immortalization-upregulated protein /DB_XRI NM_022125
 thrombospondin 1
 hypothetical protein FLJ13110
 Consensus includes gb:X69383 /DEF=H sapiens gene for T cell receptor gamma V region 5 /FEA=CDS_2 /DB_XREF=gi:37009 /UG=Hs.247801 H.sapiens gene for T cell r X69383
 sush-repeat-containing protein, X chromosome
 gb:NM_005179.1 /DEF=Homo sapiens avian lymphoma virus-deri /PROD=avian lymphoma virus-deri
 ADP-ribosylation factor 4-like
 Consensus includes gb:BF109941 /FEA=EST /DB_XREF=gi:10939631 /DB_XREF=est:7172005.x1 /CLONE=IMAGE:3527024 /UG=Hs.27183 Homo sapiens clone 23940 m BF109941
 Rho guanine exchange factor (GEF) 15
 cadherin 5, type 2, VE-cadherin (vascular epithelium)
 CD33 antigen (gp67)
 hypothetical protein FLJ13385
 cholinergic receptor, nicotinic, alpha polypeptide 3
 KIA00665 gene product

RAS protein activator like 2 NM_004841 1.41
 Consensus includes gb:AK024081.1 /DEF=Homo sapiens cDNA FLJ14019 fs, clone HEMBA1002503, /FEA=mRNA /DB_XREF=gi:10436370 /UG=Hs.287613 Homo sapie
 KIA0768 protein R50822 1.41
 Consensus includes gb:AL137312.1 /DEF=Homo sapiens mRNA; cDNA DKFZp781K23121 (from clone DKFZp781K23121), /FEA=mRNA /DB_XREF=gi:6807797 /UG=Hs. AL137312 1.41
 serine/threonine kinase 3 (STE20 homolog, yeast) NM_006281 1.41
 ventral anterior homeobox 2 NM_012476 1.41
 gb:NM_022155.1 /DEF=Homo sapiens PP3227 protein (PP3227), mRNA, /FEA=mRNA /GEN=PP3227 /PROD=PP3227 protein /DB_XREF=gi:11545901 /UG=Hs.302027 F NM_022155 1.41
 Integrin, beta 8 BC002630 1.41
 galanin AL556409 1.41
 CASP8 and FADD-like apoptosis regulator AF015452 1.41
 hypothetical protein FLJ11168 NM_018347 1.41
 carcinoembryonic antigen-related cell adhesion molecule 5 NM_004363 1.41
 DKFZP584C103 protein NM_015654 1.41
 Consensus includes gb:AV704232 /FEA=EST /DB_XREF=gi:10721550 /DB_XREF=est:AV704232 /CLONE=ADDBLB11 /UG=Hs.228096 Homo sapiens cDNA FLJ11682 fr. AV704232 1.41
 hypothetical protein NM_018819 1.41
 phosphoserine phosphatase NM_004577 1.41
 hepatocyte nuclear factor 4, gamma NM_004133 1.41
 collagen, type II, alpha 1 (primary osteoarthritis, spondyloepiphyseal dysplasia, congenital) X16468 1.41
 matrix metalloproteinase 28 NM_024302 1.41
 tripartite motif-containing 14 BC006333 1.41
 hypothetical protein MGC2652 NM_024328 1.41
 fasciculation and elongation protein zeta 1 (zyglin I) NM_005103 1.41
 hypothetical protein FKSG25 NM_022120 1.41
 low density lipoprotein receptor-related protein 4 AA584287 1.41
 B-cell CLL/lymphoma 2 NM_000657 1.40
 CGI-109 protein BG286537 1.40
 retinoid X receptor, beta BF337038 1.40
 KIAA0923 protein NM_014021 1.40
 hypothetical protein FLJ10986 NM_018291 1.40
 B9 protein BC002944 1.40
 hypothetical protein FLJ14146 NM_024709 1.40
 GDP-ucose transporter 1 NM_018389 1.40
 Consensus includes gb:AK024129.1 /DEF=Homo sapiens cDNA FLJ14067 fs, clone HEMBB1001315, /FEA=mRNA /DB_XREF=gi:10436434 /UG=Hs.287619 Homo sapie AK024129 1.40
 hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 1 NM_000862 1.40
 Consensus includes gb:BE378994 /FEA=EST /DB_XREF=gi:9324359 /DB_XREF=est:601236247f1 /CLONE=IMAGE:3608626 /UG=Hs.271717 ESTs, Weakly similar to AI BE378994 1.40
 arginine vasopressin (neurophysin II, antidiuretic hormone, diabetes insipidus, neurohypophyseal) NM_000490 1.40
 Consensus includes gb:X96937.1 /DEF=H.sapiens mRNA sequence (16p11.2), /FEA=mRNA /DB_XREF=gi:29573 /UG=Hs.307208 H.sapiens mRNA sequence (16p11.2) X96937 1.40
 met proto-oncogene (hepatocyte growth factor receptor) BE870509 1.40
 distal-less homeo box 2 BE792224 1.40
 similar to rat nuclear ubiquitous casein kinase 2 NM_022731 1.40
 dihydropyrimidinase-like 4 NM_022731 1.40
 hypothetical protein FLJ20033 AI857455 1.40
 isolated 1,4,5-trisphosphate 3-kinase C NM_017629 1.40
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 D38169 1.40
 somatostatin receptor-interacting protein AI831675 1.40
 brain specific protein NM_016148 1.40
 KIAA0769 gene product NM_016140 1.40
 PR domain containing 11 NM_014653 1.40
 hypothetical protein FLJ20510 NM_020229 1.40
 Consensus includes gb:U66046.1 /DEF=Homo sapiens clone 161455 breast expressed mRNA from chromosome X, /FEA=mRNA /DB_XREF=gi:1519271 /UG=Hs.92663 F U66046 1.40
 Jun dimerization protein p21SNF1 NM_018664 1.40
 hypothetical protein FLJ21148 NM_024860 1.40
 transducin beta-like 1 protein AF332222 1.40
 p21(CDKN1A)-activated kinase 4 AB032968 1.40
 hypothetical protein FLJ22004 NM_023161 1.40
 Consensus includes gb:BF508244 /FEA=EST /DB_XREF=gi:11591542 /DB_XREF=est:U1-H-B14-aqa-b-02-O-U1.s1 /CLONE=IMAGE:3089210 /UG=Hs.300832 ESTs, Highly BF508244 1.40
 discoidin domain receptor family, member 2 NM_006182 1.40
 Consensus includes gb:AK024525.1 /DEF=Homo sapiens cDNA: FLJ20872 fs, clone ADKA02604, /FEA=mRNA /DB_XREF=gi:10436827 /UG=Hs.306683 Homo sapiens c AK024525 1.40
 neutralized-like (Drosophila) AF028728 1.40
 tryptase, alpha AF206665 1.40
 trans-golgi network protein 2 BF081845 1.40
 epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian) AF125253 1.40
 immunoglobulin superfamily, member 6 NM_005849 1.40

Consensus includes gb:AF005082.1 /DEF=Homo sapiens skin-specific protein (xp33) mRNA, partial cds. /FEA=mRNA /GEN=xp33 /PROD=skin-specific protein /DB_XREF=AF005082
 suc1-associated neurotrophic factor target (FGFR signalling adaptor) NM_006654 1.40
 prostate tumor over expressed gene 1 U79287 1.40
 Consensus includes gb:AW503390 /FEA=EST /DB_XREF=gi:7118730 /DB_XREF=est:UH-F-BNO-akw-c02-0.U1.1 /CLONE=IMAGE:3078507 /UG=Hs.135258 ESTs. Was AW503390 1.40
 Consensus includes gb:AL137435.1 /DEF=Homo sapiens mRNA; cDNA DKFZp781G0924 (from clone DKFZp781G0924). /FEA=mRNA /DB_XREF=gi:8607994 /UG=Hs.27 AL137435 1.40
 tenascin R (restridin, jensin) NM_003285 1.40
 polymerase (RNA) II (DNA directed) polypeptide J (13.3kD) BE676209 1.40
 glycine amidinotransferase (L-arginine:glycine amidinotransferase) X88401 1.40
 KIAA1039 protein AK000478 1.40
 taste receptor, type 2, member 8 NM_023918 1.40
 gelsolin (amyloidosis, Finnish type) BE675337 1.40
 f-box and leucine-rich repeat protein 6 NM_024555 1.40
 armadillo repeat protein ALEX2 NM_014782 1.40
 hypothetical protein FLJ10851 NM_018245 1.40
 Consensus includes gb:AK023909.1 /DEF=Homo sapiens cDNA FLJ13847 fs, clone THYRO1000852, highly similar to Human branched-chain amino acid aminotransferas AK023909 1.40
 hypothetical protein FLJ22843 NM_025184 1.40
 hypothetical protein FLJ10620 NM_018157 1.40
 mitogen-activated protein kinase kinase 5 D84478 1.40
 g20 protein NM_016210 1.40
 ATPase, Class V, type 10C AB011136 1.40
 Consensus includes gb:A143878 /FEA=EST /DB_XREF=gi:3665688 /DB_XREF=est:qe04f11 x1 /CLONE=IMAGE:1738029 /UG=Hs.128291 phosphodiesterase 10A /FL=g A143878 1.40
 type 1 tumor necrosis factor receptor shedding aminopeptidase regulator AB011097 1.40
 dual specificity phosphatase 9 NM_001395 1.40
 hypothetical protein FLJ00002 AK000002 1.40
 hypothetical protein FLJ21870 NM_023016 1.40
 DKFZP434J154 protein AB011538 1.40
 slit homolog 3 (Drosophila) NM_001463 1.40
 frizzled-related protein NM_019105 1.40
 tenascin XB NM_014395 1.40
 dual adaptor of phosphotyrosine and 3-phosphoserines NM_058622 1.40
 Wiskott-Aldrich syndrome protein interacting protein AW058622 1.40
 SKI-like X15217 1.40
 spermidine/spermine N1-acetyltransferase M55580 1.40
 Machado-Joseph disease (spinocerebellar ataxia 3, autosomal dominant, ataxin 3) NM_004993 1.40
 phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 dunce homolog, Drosophila) AF012074 1.40
 general transcription factor IIIC, polypeptide 1 (alpha subunit, 220kD) U02818 1.40
 chromosome 17 open reading frame 1A U43383 1.40
 transmembrane protease, serine 2 AF270487 1.40
 RuvB-like 1 (E. coli) M34421 1.40
 Consensus includes gb:AL583528 /FEA=EST /DB_XREF=gi:12952578 /DB_XREF=est:AL583528 /CLONE=CS000004YH21 (5 prime) /UG=Hs.42915 ARP2 (actin-related eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein) AL583528 1.40
 TUJ1231-TV protein NM_016575 1.40
 secreted protein of unknown function AA679297 1.40
 lymphocyte-specific protein 1 NM_002339 1.40
 FBJ murine osteosarcoma viral oncogene homolog B NM_006732 1.40
 kinesin family member 13B NM_015254 1.40
 leukotriene C4 synthase NM_000897 1.40
 transketolase-like 1 X91817 1.40
 hypothetical protein FLJ12552 AK024318 1.40
 Fc fragment of IgG binding protein NM_003890 1.40
 testis specific protein, Y-linked M84893 1.40
 complement component 1, q subcomponent binding protein M89039 1.40
 retinoic acid induced 2 NM_021785 1.40
 Consensus includes gb:AL110206.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566N2022 (from clone DKFZp566N2022). /FEA=mRNA /DB_XREF=gi:5817125 /UG=Hs.30 AL110206 1.40
 eukaryotic translation initiation factor 3, subunit 1 (alpha, 35kD) NM_003758 1.40
 protodactherin 18 BF222893 1.40
 hypothetical protein NM_013386 1.40
 son of sevenless (Drosophila) homolog 2 L20686 1.40
 Consensus includes gb:AW168915 /FEA=EST /DB_XREF=gi:6400440 /DB_XREF=est:j15b10.x1 /CLONE=IMAGE:2657275 /UG=Hs.283946 Homo sapiens prostate-spec AW168915 1.40
 Consensus includes gb:AA156240 /FEA=EST /DB_XREF=gi:1727858 /DB_XREF=est:z50c12.x1 /CLONE=IMAGE:505366 /UG=Hs.286660 Homo sapiens cDNA: FLJ2218 AA156240 1.40
 olfactory receptor, family 5, subfamily V member 1 NM_030876 1.40
 hypothetical protein BF055311 1.40
 nucleosomal binding protein 1 NM_030763 1.40

ras-like protein
interleukin 2
sodium channel, voltage-gated, type X, alpha polypeptide
A kinase (PRKA) anchor protein (gravin) 12
tumor necrosis factor receptor superfamily, member 11a, activator of NFKB
actin A receptor, type IB
serum-inducible kinase
KIAA0643 protein
T cell receptor beta variable 17
chromosome 18 open reading frame 5
Consensus includes gb:AK000156.1 /DEF=Homo sapiens cDNA FLJ20149 fls, clone COL08213. /FEA=mRNA /DB_XREF=gi:7020059 /UG=Hs.272193 Homo sapiens cDNA AK000156
H2B histone family, member E
hypothetical protein FLJ20284
endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2
gb:NM_014823.1 /DEF=Homo sapiens KIAA0344 gene product (KIAA0344), mRNA. /FEA=mRNA /GEN=KIAA0344 /PROD=KIAA0344 gene product /DB_XREF=gi:766208 /NM_014823
SH3-domain binding protein 2
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3
gb:NM_018123.1 /DEF=Homo sapiens hypothetical protein FLJ10517 (FLJ10517), mRNA. /FEA=mRNA /GEN=FLJ10517 /PROD=hypothetical protein FLJ10517 /DB_XREF=gi:181233
zinc finger protein 8 (CAMPX1)
formyl peptide receptor-like 2
epithelial protein lost in neoplasm beta
nuclear factor /B
Consensus includes gb:A340483 /FEA=EST /DB_XREF=gi:4083689 /DB_XREF=est:qp51902.x1 /CLONE=IMAGE:1926506 /UG=Hs.132807 Homo sapiens (clone 3.8-1) A340483
Consensus includes gb:AF070579.1 /DEF=Homo sapiens clone 24487 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387951 /UG=Hs.263819 Homo sapiens clone 24487 AF070579
chromosome 1 open reading frame 25
mannosidase, alpha, class 1A, member 2
small EDRK-ichi factor 18 (centromeric)
Consensus includes gb:BE541042 /FEA=EST /DB_XREF=gi:3765786 /DB_XREF=est:801064390f.1 /CLONE=IMAGE:3450599 /UG=Hs.23240 Homo sapiens cDNA FLJ13
Hic-like, expressed in neurons (Drosophila)
HMP19 protein
hypothetical protein
Consensus includes gb:AK024108.1 /DEF=Homo sapiens cDNA FLJ14046 fls, clone HEMBA1006461. /FEA=mRNA /DB_XREF=gi:10436406 /UG=Hs.142677 Homo sapiens AK024108
tripartite motif-containing 33
protein kinase C and casein kinase substrate in neurons 3
retinal pigment epithelium-derived rodopsin homolog
carboxylesterase 3 (brain)
TPA regulated locus
Consensus includes gb:AW072461 /FEA=EST /DB_XREF=gi:6027459 /DB_XREF=est:va06c09.x1 /CLONE=IMAGE:2567536 /UG=Hs.169284 Homo sapiens phosphoribosyl-ATP synthase
zyxin
gap junction protein, beta 3, 31kD (connexin 31)
microchidia homolog (mouse)
spleen focus forming virus (SFFV) proviral integration oncogene spi.1
Consensus includes gb:AQ275374 /DEF=Homo sapiens partial GYH3 gene for immunoglobulin heavy chain V region, case 1, clone 19 /FEA=CDS /DB_XREF=gi:7573008 /AJ275374
tumor endothelial marker 5 precursor
androgen-induced prostate proliferative shutoff associated protein
protein kinase C, gamma
Consensus includes gb:AW972359 /FEA=EST /DB_XREF=gi:8162205 /DB_XREF=est:EST384450 /UG=Hs.293334 ESTs
carbonic anhydrase VIII
KIA0964 protein
guanine nucleotide binding protein (G protein), alpha 11 (Gq class)
caspase recruitment domain protein 10
KIAA1026 protein
Consensus includes gb:AU144277 /FEA=EST /DB_XREF=gi:11005798 /DB_XREF=est:AU144277 /CLONE=HEMBA1001415 /UG=Hs.296525 Homo sapiens cDNA FLJ101
cold shock domain protein A
angiotensin receptor 2
matrix metalloproteinase 24 (membrane-inserted)
aryl hydrocarbon receptor nuclear translocator 2
gb:NM_018628.1 /DEF=Homo sapiens hypothetical protein PRO2472 (PRO2472), mRNA. /FEA=mRNA /GEN=PRO2472 /PROD=hypothetical protein PRO2472 /DB_XREF=gi:18628
hypothetical protein FLJ23239
hypothetical protein FLJ10770
vasoactive intestinal peptide receptor 2
hypothetical protein FLJ21007
paraneoplastic antigen MA2

sialic acid binding Ig-like lectin 6	D88359	1.39
cytochrome, alpha	NM_001392	1.39
peroxisomal trans 2-oxoyl CoA reductase; putative short chain alcohol dehydrogenase	NM_018441	1.39
Sjogren syndrome antigen B (autoantigen La)	BC532828	1.39
RFamide-related peptide precursor	NM_022150	1.39
dual specificity phosphatase 3 (vaccinia virus phosphatase VH11-related)	BC002882	1.39
nuclear receptor coactivator 1	U18179	1.39
hypothetical protein FLJ20753	NM_017950	1.39
homeo box D13	NM_000523	1.39
Consensus includes gb:BG251521 /FEA=EST /DB_XREF=gi:127181337 /DB_XREF=est:602363985f1 /CLONE=IMAGE:4472160 /UG=Hs.16193 Homo sapiens mRNA; cdi	BC251521	1.39
hypothetical protein FLJ10803	AJ554945	1.39
hypothetical protein FLJ20761	NM_017954	1.39
CGI-12 protein	NM_015942	1.39
spinocerebellar ataxia 8	AF126749	1.39
hypothetical protein FLJ12168	NM_024682	1.39
Consensus includes gb:AK022999.1 /DEF=Homo sapiens cDNA FLJ12937 fls, clone NT2R2P005020. /FEA=mRNA /DB_XREF=gi:10434712 /UG=Hs.14846 Homo sapiens	AK022999	1.39
fibroblast growth factor 1 (ecdic)	X59085	1.39
member of MYST family histone acetyl transferases, homolog of Drosophila MOF	AL050395	1.39
protein tyrosine phosphatase, receptor type, G	NM_002841	1.39
gastrulation brain homeo box 1	L11239	1.39
glioma-associated oncogene homolog (zinc finger protein)	NM_005269	1.39
cadherin 11, type 2, OB-cadherin (osteoblast)	D21254	1.39
neuronal PAS domain protein 2	U77970	1.39
hypothetical protein FLJ20513	NM_017855	1.39
signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	NM_005643	1.39
Consensus includes gb:BE673151 /FEA=EST /DB_XREF=gi:10033692 /DB_XREF=est:7d30a1.1.x1 /CLONE=IMAGE:3248732 /UG=Hs.91626 Homo sapiens clone B18 un	BE673151	1.39
membrane-bound transcription factor protease, site 1	BE890314	1.39
Consensus includes gb:AL530331 /FEA=EST /DB_XREF=gi:12793824 /DB_XREF=est:AL530331 /CLONE=CS0DD007YMO6 (3 prime) /UG=Hs.111243 MADS box transcri	AL530331	1.39
colon cancer-associated protein Mtc1	NM_013326	1.39
early growth response 4	NM_001965	1.39
PH domain containing protein in retina 1	AF001583	1.39
collagen, type X, alpha 1 (Schmid metaphyseal chondrodysplasia)	A1376003	1.39
Consensus includes gb:AK021741.1 /DEF=Homo sapiens cDNA FLJ11679 fls, clone HEMBA1004807. /FEA=mRNA /DB_XREF=gi:10432986 /UG=Hs.287439 Homo sapie	AK021741	1.39
peptidylprolyl isomerase E (cyclophilin E)	AF104013	1.39
KIAA0560 gene product	AB011132	1.39
calpain 3, (p94)	AF127764	1.39
pyruvate dehydrogenase kinase, isoenzyme 3	NM_005391	1.39
fibronectin leucine rich transmembrane protein 2	NM_013231	1.39
matrix metalloproteinase 16 (membrane-inserted)	AB009303	1.39
hypothetical protein FLJ22596	NM_025086	1.39
myosin VI	AA877789	1.39
musashi homolog 1 (Drosophila)	AC003982	1.39
heat shock transcription factor 1	NM_005526	1.39
CGI-142	AL133102	1.39
hypothetical protein MGC3103	NM_024036	1.39
hemoglobin, alpha 2	T50399	1.39
solute carrier family 12, (potassium/chloride transporter) member 5	AF208159	1.39
gb:AF130093.1 /DEF=Homo sapiens clone FLB9835 PRO2660 mRNA, complete cds. /FEA=mRNA /PROD=PRO2660 /DB_XREF=gi:11493490 /UG=Hs.302153 Homo sap	AF130093	1.39
S100 calcium binding protein A3	NM_002960	1.39
platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog)	NM_002608	1.39
Consensus includes gb:BF446180 /FEA=EST /DB_XREF=gi:11511318 /DB_XREF=est:7p33e08.x1 /CLONE=IMAGE:3647703 /UG=Hs.41639 programmed cell death 2	BF446180	1.39
KIAA0573 protein	A827677	1.39
hypothetical protein FLJ11756	NM_024606	1.39
myeloid/lymphoid or mixed-lineage leukemia (rithorax homolog, Drosophila); translocated to, 4	AB016898	1.39
pvt-1 (murine) oncogene homolog, MYC activator	M34428	1.39
four jointed box 1 (Drosophila)	NM_014344	1.39
calmodulin 3 (phosphorylase kinase, delta)	NM_005184	1.39
fibronectin leucine rich transmembrane protein 1	AF169875	1.39
keratin associated protein 1.1	NM_030967	1.39
matrix metalloproteinase 24 (membrane-inserted)	AL121753	1.39
protein kinase, cGMP-dependent, type I	D45864	1.39
AtPase, Class VI, type 11A	BF439472	1.39
Consensus includes gb:N54942 /FEA=EST /DB_XREF=gi:1198282 /DB_XREF=est:y38008.s1 /CLONE=IMAGE:244983 /UG=Hs.276590 ESTs	N54942	1.39

T-box 4	NM_018488	1.39
KIAA1614 protein	AB048834	1.39
pro-melanin-concentrating hormone-like 1	S64288	1.39
zinc finger protein 161	AU146275	1.39
epidermal growth factor receptor substrate EPS1R	AV710549	1.39
Consensus includes gb:AU145357 /FEA=EST /DB_XREF=est:AU145357 /CLONE=HEMBA1004611 /UG=Hs.296700 Homo sapiens cDNA FLJ12	AU145357	1.39
hypothetical protein FLJ20624	NM_017906	1.39
collagen, type I, alpha 2	AA788711	1.39
RAN binding protein 7	BG291787	1.39
KIAA0754 protein	AW663712	1.39
Consensus includes gb:AL049263.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F133 (from clone DKFZp564F133) /FEA=mRNA /DB_XREF=gi:4500011 /UG=Hs.30821	AL049263	1.39
hypothetical protein FLJ21125	NM_024627	1.39
thymosin, beta 10	AF060913	1.39
lucosyltransferase 4 (alpha 1(3) lucosyltransferase, myeloid-specific)	M58596	1.39
Consensus includes gb:AL050328 /DEF=Human DNA sequence from clone RP11-145L22 on chromosome 6p21.32-22.2. Contains the gene for myelinoligodendrocyte gly	AL050328	1.39
catenin (cadherin-associated protein), alpha pseudogene 1	Z37894	1.39
coagulation factor II (thrombin) receptor-like 1	NM_005242	1.39
DKFZP564J0206 protein	A927000	1.39
glutamate receptor, ionotropic, kainate 3	NM_000831	1.39
monoamine oxidase A	NM_000240	1.39
kalikrein 7 (chymotryptic, stratum corneum)	NM_005046	1.39
KIAA0902 protein	NM_014927	1.39
hepatocyte growth factor (hepatopolein A; scatter factor)	M77227	1.39
GRB2-related adaptor protein	NM_006613	1.39
EGF-containing fibulin-like extracellular matrix protein 2	AB030855	1.39
hypothetical protein	NM_018985	1.39
cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 1	NM_000499	1.39
DnaJ (Hsp40) homolog, subfamily B, member 6	AL353379	1.39
Consensus includes gb:BC396614 /FEA=EST /DB_XREF=gi:13290062 /DB_XREF=est:602459468F1 /CLONE=IMAGE:4581928 /UG=Hs.154999 ESTs, Moderately similar	BG396614	1.39
retin (Reed-Stenberg cell-expressed intermediate filament-associated protein)	NM_002566	1.39
transmembrane protein 2	NM_013390	1.39
glyche-N-acyltransferase	NM_005838	1.39
protein-kinase, interferon-inducible double stranded RNA dependent inhibitor, repressor of (PS8 repressor)	AF081567	1.39
basement membrane-induced gene	NM_004848	1.39
hypothetical protein MGC5395	NM_024060	1.39
coronin, actin binding protein, 2B	BF639649	1.39
acidic epididymal glycoprotein-like 1	NM_001131	1.39
Consensus includes gb:AI378979 /FEA=EST /DB_XREF=gi:4188832 /DB_XREF=est:td0607.x1 /CLONE=IMAGE:2067085 /UG=Hs.292911 ESTs	AI378979	1.39
EphB1	AF037334	1.39
tubulin, gamma 2	NM_016437	1.39
hypothetical protein FLJ12891	NM_024950	1.39
spondin 1, (f-spondin) extracellular matrix protein	A1855290	1.39
Consensus includes gb:Y14488.1 /DEF=Homo sapiens mRNA for putative 14kD protein containing SHMT homology, clone pUS1215. /FEA=mRNA /PROD=14 kDa protein	Y14488	1.39
PTEN induced putative kinase 1	AF316873	1.39
PTEN induced putative kinase 1	BF432478	1.39
SMA3	NM_006780	1.39
acyl-Coenzyme A dehydrogenase, C-2 to C-3 short chain	NM_000017	1.39
H2A histone family, member X	BE901081	1.39
calpain 6	AJ003388	1.39
hypothetical protein FLJ11126	BC005162	1.39
special AT-rich sequence binding protein 1 (binds to nuclear matrix/scaffold-associating DNA's)	NM_002971	1.39
6-pyruvoyl-tetrahydropterin synthase/dimerization cofactor of hepatocyte nuclear factor 1 alpha (TCF-1)	NM_000281	1.39
Consensus includes gb:AL050145.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586C2020 (from clone DKFZp586C2020) /FEA=mRNA /DB_XREF=gi:4894356 /UG=Hs.22	AL050145	1.39
RAB11B, member RAS oncogene family	AL575337	1.39
vaccinia related kinase 1	NM_003384	1.39
glutaminase	AF097493	1.39
dihydropyrimidinase-like 4	NM_008426	1.39
BAI1-associated protein 3	BE328802	1.39
eukaryotic translation initiation factor 2C, 1	NM_012189	1.39
KIAA0632 protein	AK001815	1.39
polyadenylate binding protein-interacting protein 1	AL136920	1.39
testis specific protein, Y-linked	NM_003308	1.39
like-glycosyltransferase	BF057493	1.39

collagen, type IV, alpha 6
Consensus includes gb:AL117571.1 /DEF=Homo sapiens mRNA; cDNA DKFZ564O1172 (from clone DKFZ564O1172). /FEA=mrna /DB_XREF=gi:5912123 /UG=Hs.24
1.39

zeta-chain (TCR) associated protein kinase (70 kD)
Consensus includes gb:AA628440 /FEA=EST /DB_XREF=gi:2540827 /DB_XREF=est:af2802.s1 /CLONE=IMAGE:1032795 /UG=Hs.83623 nuclear receptor subfamily 1, gi:AA628440
1.39

phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 duncle homolog, Drosophila)
Consensus includes gb:AL121975 /FEA=Human DNA sequence from clone RP3-422B11 on chromosome 6p11.2-12.3 Contains ESTs, STSs and GSSs. Contains part of the AL121975
1.39

zinc finger protein
Consensus includes gb:AA349848 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

plexin A1
Consensus includes gb:BE619220 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

KIAA0871 protein
Consensus includes gb:AA861784 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

peptidylprolyl isomerase C (cyclophilin C)
Consensus includes gb:AA628440 /FEA=EST /DB_XREF=gi:2540827 /DB_XREF=est:af2802.s1 /CLONE=IMAGE:1032795 /UG=Hs.83623 nuclear receptor subfamily 1, gi:AA628440
1.39

Consensus includes gb:AA628440 /FEA=EST /DB_XREF=gi:2540827 /DB_XREF=est:af2802.s1 /CLONE=IMAGE:1032795 /UG=Hs.83623 nuclear receptor subfamily 1, gi:AA628440
1.39

Consensus includes gb:AL121975 /FEA=Human DNA sequence from clone RP3-422B11 on chromosome 6p11.2-12.3 Contains ESTs, STSs and GSSs. Contains part of the AL121975
1.39

acid phosphatase 5, tartrate resistant
Consensus includes gb:AA016111 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

myosin XVA
Consensus includes gb:NN_016239 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

stratulin
Consensus includes gb:NN_016239 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

2,3-bisphosphoglycerate mutase
Consensus includes gb:NN_016239 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

tumor protein 63 kDa with strong homology to p53
Consensus includes gb:NN_016239 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

acetyl-Coenzyme A transporter
Consensus includes gb:NN_016239 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

hypothetical protein FLJ14803
Consensus includes gb:NN_016239 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

epilepsy, progressive myoclonus type 2, Lafora disease (laforin)
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

hypothetical protein FLJ13352
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

Thy-1 cell surface antigen
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

hypothetical protein FLJ10178
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

regulator of G-protein signalling 12
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

calcium channel, voltage-dependent, alpha 1I subunit
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

heart and neural crest derivatives expressed 2
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

vanilloid receptor subtype 1
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

KIAA1822 protein
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

Rac GTPase activating protein 1
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

tripartite motif-containing 31
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

KIAA0685 gene product
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

SH3 domain binding glutamic acid-rich protein
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

guanylate binding protein 1, interferon-inducible, 67kD
Consensus includes gb:BF940043 /FEA=EST /DB_XREF=gi:12357363 /DB_XREF=est:nac681f2.x1 /CLONE=IMAGE:3439271 /UG=Hs.62041 nidogen (enactin) /FL=gb:MC BF940043
1.39

Consensus includes gb:BE877397 /FEA=EST /DB_XREF=gi:10326173 /DB_XREF=est:1801485434F1 /CLONE=IMAGE:3887938 /UG=Hs.240140 Human DNA sequence fr
1.39

hypothetical protein
Consensus includes gb:BE877397 /FEA=EST /DB_XREF=gi:10326173 /DB_XREF=est:1801485434F1 /CLONE=IMAGE:3887938 /UG=Hs.240140 Human DNA sequence fr
1.39

MAC, mothers against decapentaplegic homolog 5 (Drosophila)
Consensus includes gb:BE877397 /FEA=EST /DB_XREF=gi:10326173 /DB_XREF=est:1801485434F1 /CLONE=IMAGE:3887938 /UG=Hs.240140 Human DNA sequence fr
1.39

hematopoietic PBX-interacting protein
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mrna /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

extracellular matrix protein 2, female organ and adipocyte specific
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mrna /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

WD repeat domain 7
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mrna /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

hypothetical protein FLJ21477
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mrna /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mrna /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

CGI-14 protein
Consensus includes gb:U79295.1 /DEF=Human clone 23861 mRNA sequence. /FEA=mrna /DB_XREF=gi:1710277 /UG=Hs.90866 Human clone 23861 mRNA sequence
1.39

tetratropopeptide repeat domain 3
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

vitamin D (1,25-dihydroxyvitamin D3) receptor
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

wingless-type MMTV integration site family, member 3
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

laminin, beta 1
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

neurogenin 2
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

collagen, type IV, alpha 5 (Alport syndrome)
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

nucleolar protein 3 (epitopsin repressor with CARD domain)
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

CD8 antigen
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

CD32 antigen, zeta polypeptide (TIT3 complex)
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

arginine vasopressin receptor 1B
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

vanilloid receptor-related osmotically activated channel; OTRPC4 protein
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

hypothetical protein MGC4293
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

pre-B-cell leukemia transcription factor 2
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA FLJ12301 fls. clone MAMMA1001858. /FEA=mrna /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie
1.38

Consensus includes gb:AL583520 /FEA=EST /DB_XREF=gi:12952562 /DB_XREF=est:AL583520 /CLONE=CSDDC024YE13 (5 prime) /UG=Hs.182183 Homo sapiens mRi
1.38

cell division cycle 27
Consensus includes gb:AL583520 /FEA=EST /DB_XREF=gi:12952562 /DB_XREF=est:AL583520 /CLONE=CSDDC024YE13 (5 prime) /UG=Hs.182183 Homo sapiens mRi
1.38

transcobalamin II; macrocytic anemia
Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA; FLJ21342 fls. clone COL02873. /FEA=mrna /DB_XREF=gi:10437428 /UG=Hs.306739 Homo sapiens cC
1.38

Consensus includes gb:AK022363.1 /DEF=Homo sapiens cDNA; FLJ21342 fls. clone COL02873. /FEA=mrna /DB_XREF=gi:10437428 /UG=Hs.306739 Homo sapiens cC
1.38

Consensus includes gb:U38460 /DEF=Human RNA binding motif (RBM) gene, partial cds /FEA=CDS /DB_XREF=gi:1061395 /UG=Hs.326802 Human RNA binding motif (F U38460
1.38

heterogeneous nuclear ribonucleoprotein H3 (2H9) 1.38
translocase of inner mitochondrial membrane 8 homolog A (yeast) 1.38
double homeobox_3 1.38
annexin A6 1.38
Consensus includes gb:AK021633.1 /DEF=Homo sapiens cDNA FLJ11571 fs, clone HEMBA1003328 /FEA=mRNA /DB_XREF=gi: 0432853 /UG=Hs.306808 Homo sapiens 1.38
melanocortin 4 receptor 1.38
hypothetical protein FLJ23033 1.38
solute carrier family 25 (mitochondrial carrier; Graves disease autoantigen), member 16 1.38
similar to DNA-directed RNA polymerase I (135 kDa) 1.38
X-prolyl aminopeptidase (aminopeptidase P) 1, soluble 1.38
syndecan 1 1.38
protein (peptidyl-prolyl cis/trans isomerase) NIMA-interacting 1-like 1.38
ret finger protein-like 2 1.38
hypothetical protein PRO1230 1.38
glutamate receptor, metabotropic 1 1.38
hypothetical protein FLJ22087 1.38
estrogen receptor 2 (ER beta) 1.38
ATP-binding cassette, sub-family B (MDR/TAP), member 6 1.38
nucleophosmin (nucleolar phosphoprotein B23, numatrin) 1.38
protease, serine, 7 (enterokinase) 1.38
hypothetical protein MGC10848 1.38
KIAA0810 protein 1.38
small proline-rich protein 1A 1.38
Consensus includes gb:AK026273.1 /DEF=Homo sapiens cDNA: FLJ22620 fs, clone HS05629 /FEA=mRNA /DB_XREF=gi: 10439077 /UG=Hs.306834 Homo sapiens cDI AK026273 1.38
Consensus includes gb:AL049369.1 /DEF=Homo sapiens mRNA, cDNA DKFZp586D0518 (from clone DKFZp586D0518) /FEA=mRNA /DB_XREF=gi: 4500160 /UG=Hs.25 AL049369 1.38
chromosome 19 open reading frame 7 1.38
lethal (3) malignant brain tumor l(3)mb1 protein (Drosophila) homolog 1.38
KIAA1052 protein 1.38
Consensus includes gb:AL117549.1 /DEF=Homo sapiens mRNA, cDNA DKFZp586H0123 (from clone DKFZp586H0123) /FEA=mRNA /DB_XREF=gi: 5912083 /UG=Hs.26 AL117549 1.38
immunoglobulin lambda locus 1.38
protocadherin 17 1.38
hypothetical protein FLJ21916 1.38
solute carrier family 2 (facilitated glucose transporter), member 3 1.38
thioredoxin reductase beta 1.38
KIAA0143 protein 1.38
dolichyl-phosphate (UDP-N-acetylglucosamine) N-acetylglucosaminylphosphotransferase 1 (GlcNAc-1-P transferase) 1.38
Consensus includes gb:AL080107.1 /DEF=Homo sapiens mRNA, cDNA DKFZp586D0522 (from clone DKFZp586D0522) /FEA=mRNA /DB_XREF=gi: 5262532 /UG=Hs.16 AL080107 1.38
ADP-ribosylation factor-like 7 1.38
hypothetical protein PP3501 1.38
pancreatic lipase-related protein 1 1.38
hypothetical protein PRO2521 1.38
CCR4-NOT transcription complex, subunit 8 1.38
myeloid/lymphoid or mixed-lineage leukemia (rithorax homolog, Drosophila); translocated to, 10 1.38
B lymphoid tyrosine kinase 1.38
bromodomain-containing 3 1.38
fatty acid hydroxylase 1.38
carboxylate kinase-like 1.38
M24537B subtilis phoB, phoA genes corresponding to nucleotides 2017-3334 of M24537 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively) 1.38
KIAA1751 protein 1.38
solute carrier family 21 (prostaglandin transporter), member 2 1.38
N-acetylglucosamine-phosphate mutase 1.38
DKFZP5841122 protein 1.38
NAD(P)H dehydrogenase, quinone 1 1.38
inhibitor of DNA binding 4, dominant negative helix-loop-helix protein 1.38
a disintegrin and metalloproteinase domain 12 (meltrin alpha) 1.38
cadherin, EGF LAG seven-pass G-type receptor 3 (flamingo homolog, Drosophila) 1.38
maltase-glucoamylase (alpha-glucosidase) 1.38
DKFZP4348204 protein 1.38
X03453 Bacteriophage P1 cre recombinase protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively) 1.38
serine (or cysteine) proteinase inhibitor, clade H (heat shock protein 47), member 1 1.38
tumor up-regulated CARD-containing antagonist of caspase nine 1.38
filamin C, gamma (actin binding protein 280) 1.38
T-box 1 1.38

inositol 1,4,5-trisphosphate receptor, type 3
 hypothetical protein MGC10771
 neuropilin 3
 Vertebrate LIN7 homolog 1, Tax interaction protein 33
 Vertebrate LIN7 homolog 1, Tax interaction protein 2
 chloride channel, calcium activated, family member 2
 endoglin (Osler-Rendu-Weber syndrome 1)
 sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3F
 Consensus includes gb:AA160522 /FEA=EST /DB_XREF=gi:1735889 /DB_XREF=est:z46b10.s1 /CLONE=IMAGE:632731 /UG=Hs.301667 Homo sapiens mRNA; cDNA (AA160522)
 Consensus includes gb:AL080232.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586A061 (from clone DKFZp586A061) /FEA=mRNA /DB_XREF=gi:5262725 /UG=Hs.2208 /AL080232
 lms-related tyrosine kinase 3
 acid fibroblast growth factor-like protein
 keratin 1 (epidermolytic hyperkeratosis)
 neuronal potassium channel alpha subunit
 solute carrier family 4, sodium bicarbonate cotransporter, member 4
 Consensus includes gb:AW576871 /FEA=EST /DB_XREF=gi:7251920 /DB_XREF=est:RC2-BT0318-110100-012-e02 /UG=Hs.201420 ESTs, Moderately similar to ALU7_F-AW576871
 hypothetical protein PRO2958
 transducin-like enhancer of split 2 (E(spl) homolog, Drosophila)
 Consensus includes gb:AW054711 /FEA=EST /DB_XREF=gi:5920414 /DB_XREF=est:wc86g08.x1 /CLONE=IMAGE:2566718 /UG=Hs.121591 Human DNA sequence from AW054711
 M99436
 Consensus includes gb:AF003738.1 /DEF=Homo sapiens chromosome 21q22.1 anonymous mRNA sequence. /FEA=mRNA /DB_XREF=gi:2197087 /UG=Hs.268526 Homr AF003738
 ATPase, Ca⁺⁺ transporting, cardiac muscle, slow twitch 2
 Consensus includes gb:A004009 /FEA=EST /DB_XREF=gi:3213519 /DB_XREF=est:ou03410.x1 /CLONE=IMAGE:1625203 /UG=Hs.130816 ESTs, Moderately similar to A004009
 membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE, high affinity I, receptor for: beta polypeptide)
 Consensus includes gb:A1249275 /DEF=Homo sapiens partial MTHFR gene for methylenetetrahydrofolate reductase /FEA=CDS /DB_XREF=gi:7271187 /UG=Hs.214142 F-A1249275
 Consensus includes gb:A1693193 /FEA=EST /DB_XREF=gi:4970533 /DB_XREF=est:wd6802.x1 /CLONE=IMAGE:2338763 /UG=Hs.247551 metaxin 1
 N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1
 gb:AF274945.1 /DEF=Homo sapiens PNAS-20 mRNA, complete cds. /FEA=mRNA /PROD=PNAS-20 /DB_XREF=gi:12751054 /FL=gb:AF274945.1
 vacuolar protein sorting 33B (yeast)
 hypothetical protein FLJ10846
 hypothetical protein MGC2991
 hypothetical protein FLJ12750
 hypothetical protein MGC3129 similar to ganglioside-induced differentiation-associated protein
 calcium channel, voltage-dependent, gamma subunit 3
 KIAA0197 protein
 dom-3 homolog 2 (C. elegans)
 Consensus includes gb:AL110201.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586F1622 (from clone DKFZp586F1622) /FEA=mRNA /DB_XREF=gi:5917120 /UG=Hs.271 /AL110201
 U87459
 cancer/testis antigen
 double ring-finger protein, Doriin
 transforming growth factor, beta-induced, 68kD
 putative leukocyte platelet-activating factor receptor
 annexin A2 pseudogene 3
 UC28 protein
 hemoglobin, delta
 inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase complex-associated protein
 prostaglandin E receptor 3 (subtype EP3)
 a disintegrin and metalloproteinase domain 5
 sarcosin (Kras oncogene-associated gene)
 protein phosphatase 2, regulatory subunit B (B56), beta isoform
 Consensus includes gb:A1907083 /FEA=EST /DB_XREF=gi:6497611 /DB_XREF=est:PM-BT134-050499-650 /UG=Hs.124620 ESTs
 KIAA1031 protein
 B-factor, properdin
 glucagon-like peptide 1 receptor
 hypothetical protein FLJ12568
 phosphodiesterase 1C; calmodulin-dependent (70kD)
 KIAA1099 protein
 gamma-glutamyl carboxylase
 hypothetical protein DKFZp434C0328
 hypothetical protein FLJ11807
 hypothetical protein FLJ20420
 likely ortholog of mouse NPC derived proline rich protein 1
 hypothetical protein FLJ21511
 ephrin-B2
 CGI-130 protein

fibroblast growth factor 5 1.38 AF171928
outer dense fibre of sperm tails 1 1.38 NM_024410
KIA0082 protein 1.38 AL136827
centaurin, delta 2 1.38 AB018325
activin A receptor, type IB 1.38 NM_020327
cadherin 12, type 2 (N-cadherin 2) 1.38 NM_016542
serine/threonine protein kinase MASK 1.38 L33477
Consensus includes gb:51708 /FEA=EST /DB_XREF=gi:1192874 /DB_XREF=est:y72e03.s1 /CLONE=IMAGE:279100 /UG=Hs.182723 Homo sapiens mRNA from chrom N51708
heat shock 70kD protein-like 1 1.38 NM_016542
tropoin 1, skeletal, fast 1.38 D85730
Consensus includes gb:AF043584.1 /DEF=Homo sapiens clone ASMneg1-b1 Immunoglobulin lambda chain V J region, (JGL) mRNA, partial cds. /FEA=mRNA /GEN=JGL
hypothetical protein MGC3130 1.38 NM_003282
hypothetical protein FLJ12076 1.38 AF043584
cardiag oligomeric matrix protein (pseudosarcomere), epiphyseal dysplasia 1, multiple 1.38 NM_024032
hypothetical protein FLJ12345 1.38 NM_025187
Consensus includes gb:AA478678 /FEA=EST /DB_XREF=gi:2205564 /DB_XREF=est:zu43a04.s1 /CLONE=IMAGE:740718 /UG=Hs.203269 ESTs, Moderately similar to AI AA478678
hypothetical protein FLJ14280 1.38 NM_024987
gb:BC005043.1 /DEF=Homo sapiens, Similar to tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain, clone MGC:12787, mRNA, BC005043
acid sphingomyelinase-like phosphodiesterase 1.38 NM_024886
hypothetical protein FLJ11783 1.38 BC005043
selectin L (lymphocyte adhesion molecule 1) 1.38 NM_014474
major histocompatibility complex, class I-like sequence 1.38 NM_024891
Consensus includes gb:AK023958.1 /DEF=Homo sapiens cDNA FLJ13897 fs, clone THYRO1001706, /FEA=mRNA /DB_XREF=gi:10436079 /UG=Hs.301468 Homo sapiens KIA0084 protein 1.38 NM_000855
fibrillin 1 (Marfan syndrome) 1.38 AF010447
amylo-1,8-glucosidase, 4-alpha-glucanotransferase (glycogen debranching enzyme, glycogen storage disease type III) 1.38 NM_014896
golgii autoantigen, golgin subfamily b, macrogolin (with transmembrane signal), 1 1.38 NM_000138
solite carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7 1.38 NM_006845
putative tumor suppressor 1.38 N53479
complement component 1, q subcomponent, alpha polypeptide 1.38 H40895
debranching enzyme homolog 1 (S. cerevisiae) 1.38 BC004483
heat shock 27kD protein 2 1.37 NM_015991
DNA2 DNA replication helicase 2-like (yeast) 1.37 NM_016216
rhodopsin kinase 1.37 NM_001541
chromosome 4 open reading frame 5 1.37 D42046
Q antigen 6 1.37 NM_002929
likely ortholog of mouse tumor necrosis-alpha-induced adipose-related protein 1.37 AY013295
Consensus includes gb:AK022280.1 /DEF=Homo sapiens cDNA FLJ12218 fs, clone MAMMA1001075, moderately similar to Homo sapiens CGI-72 protein mRNA, /FEA=mRNA /DB_XREF=gi:10436079 /UG=Hs.225433 Homo sapiens mRNA, AL046350
hemochromatosis 1.37 NM_001476
keratin, hair, acidic, 5 1.37 AF144240
bulbous pemphigoid antigen 1 (230/240kD) 1.37 X80763
WD repeat domain 4 1.37 BC001074
Consensus includes gb:AL046350 /FEA=EST /DB_XREF=gi:5434427 /DB_XREF=est:DKFZp434J097_s1 /CLONE=DKFZp434J097 /UG=Hs.225433 Homo sapiens mRNA, AL046350
CGI-60 protein 1.37 NM_016008
myosin, heavy polypeptide 8, skeletal muscle, perinatal 1.37 M36769
Consensus includes gb:AA029811 /FEA=EST /DB_XREF=gi:1496048 /DB_XREF=est:zk07e10.s1 /CLONE=IMAGE:469866 /UG=Hs.79531 Homo sapiens TTF-I interacting AA029811
arminxin A10 1.37 AF196478
hypothetical protein MGC4093 1.37 NM_030578
Consensus includes gb:U85992.1 /DEF=Human clone IMAGE:35527 unknown protein mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1835745 /UG=Hs.t U85992
phospholipid scramblase 4 1.37 NM_020353
arginine-glutamic acid dipeptide (RE) repeats 1.37 A920976
gap junction protein, alpha 1, 43kD (connexin 43) 1.37 NM_000165
mucin and cadherin-like 1.37 NM_017717
hypothetical protein FLJ11029 1.37 NM_018304
Consensus includes gb:BG236220 /FEA=EST /DB_XREF=gi:12750151 /DB_XREF=est:maf25h11.x1 /CLONE=IMAGE:4142085 /UG=Hs.323067 Homo sapiens mRNA, cdt BG236220
GPI anchored molecule like protein 1.37 NM_002066
plakophilin 2 1.37 AA688057
hypothetical protein from BCRA2 region 1.37 U50532
gamma-aminobutyric acid (GABA) A receptor, delta 1.37 NM_000815
growth arrest and DNA-damage-inducible, beta 1.37 AV658884
aristaeless homeobox (Drosophila) 1.37 A1469991
hypothetical protein FLJ21065 1.37 NM_022481
major histocompatibility complex, class II, DO alpha 1 1.37 X00452

clickip1 homolog 1 (Xenopus laevis) NM_012242 1.37
 plakophilin 3 BF001503 1.37
 ecdonucleotide triphosphatase diphosphohydrolase 1 NM_0011776 1.37
 cholinergic receptor, nicotinic, beta polypeptide 4 NM_000750 1.37
 Consensus includes gb:AL080180.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434M091 (from clone DKFZp434M091) /FEA=mRNA /DB_XREF=gi:5282656 /UG=Hs.1822 AL080180 1.37
 transcription factor 4 NM_003189 1.37
 protein kinase C-like 2 BG252065 1.37
 dysstrobrein, alpha U46745 1.37
 neurotrophic tyrosine kinase, receptor, type 3 S76476 1.37
 KIAA1655 protein NM_0051442 1.37
 prolidase-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase, Ehlers-Danlos syndrome type VI) NM_000302 1.37
 v-maf musculoaponeurotic fibrosarcoma oncogene homolog G (avian) NM_002359 1.37
 small nuclear RNA activating complex, polypeptide 2, 45kD NM_003083 1.37
 growth hormone 1 NM_022559 1.37
 sentrin/SUMO-specific protease 3 AK000923 1.37
 nerve growth factor, beta polypeptide NM_002506 1.37
 gb:NM_019841.1 /DEF=Homo sapiens epithelial calcium channel 1 (ECAC1), mRNA, /FEA=mRNA /GEN=ECAC1 /PROD=epithelial calcium channel 1 /DB_XREF=gi:97896 NM_019841 1.37
 siruin alien mating type information regulation 2 homolog 3 (S. cerevisiae) AA92888 1.37
 hypothetical protein MGC2376 NM_023930 1.37
 BCL2-associated ataxogene 4 NM_004874 1.37
 polydenylate binding protein-interacting protein 1 AW613203 1.37
 amine oxidase, copper containing 3 (vascular adhesion protein 1) NM_003734 1.37
 cholinergic receptor, nicotinic, beta polypeptide 3 X87513 1.37
 potassium inwardly-rectifying channel, subfamily J, member 3 U39196 1.37
 collagen, type IX, alpha 3 NM_001853 1.37
 regulator of G-protein signalling 5 A183997 1.37
 KIAA0241 protein D87682 1.37
 endoglin (Osler-Rendu-Weber syndrome 1) BE732652 1.37
 transcription factor AP-4 (activating enhancer binding protein 4) NM_003223 1.37
 adenylylase cyclase 1 (brain) L05500 1.37
 neurotrophic tyrosine kinase, receptor, type 2 BF674712 1.37
 phosphatidylinositol glycan, class C NM_002642 1.37
 Consensus includes gb:AK053367 /FEA=EST /DB_XREF=gi:1544893 /DB_XREF=est:75h04.s1 /CLONE=IMAGE:364855 /UG=Hs.51515 Homo sapiens mRNA; cDNA DI AA053367 1.37
 cadherin 18, type 2 NM_004934 1.37
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 34 AA039439 1.37
 mesenchymal stem cell protein DSC43 AF242768 1.37
 protein phosphatase 1, regulatory (inhibitor) subunit 12A BE737620 1.37
 hypothetical protein NM_019016 1.37
 RAB28, member RAS oncogene family NM_014353 1.37
 Consensus includes gb:AL134453 /FEA=EST /DB_XREF=gi:6602840 /DB_XREF=est:DKFZp547J135.s1 /CLONE=IMAGE:2443339 /UG=Hs.119883 ESTs A880903 1.37
 chemokine (C-X-C) receptor 1 U20350 1.37
 hypothetical protein FLJ12810 NM_024782 1.37
 integrin, alpha V (vitronectin receptor, alpha polypeptide, antigen CD51) 2 AL134453 1.37
 ATPase, Ca++ transporting, plasma membrane 2 A093579 1.37
 CGI-116 protein M97260 1.37
 Consensus includes gb:AU158566 /FEA=EST /DB_XREF=gi:11020087 /DB_XREF=est:AU158566 /CLONE=PLACE3000322 /UG=Hs.287635 Homo sapiens cDNA FLJ143 AU158566 1.37
 calcium channel, voltage-dependent, L type, alpha 1D subunit NM_024873 1.37
 hypothetical protein FLJ21162 BE550599 1.37
 calcium channel, voltage-dependent, L type, alpha 1D subunit U34846 1.37
 aquaporin 4 U46024 1.37
 myotubular myopathy 1 NM_018280 1.37
 hypothetical protein FLJ10945 NM_015227 1.37
 chromosome 21 open reading frame 80 A1694562 1.37
 collagen, type IV, alpha 3 (Goodpasture antigen) NM_025260 1.37
 G88 protein NM_030915 1.37
 hypothetical protein DKFZp566J091 NM_021098 1.37
 calcium channel, voltage-dependent, alpha 1H subunit NM_021098 1.37
 Consensus includes gb:AK024109.1 /DEF=Homo sapiens cDNA FLJ14047.1s, clone HEMBA1006817 /FEA=mRNA /DB_XREF=gi:10436407 /UG=Hs.301186 Homo sapiens cDNA AK024109 1.37
 Consensus includes gb:AK025029.1 /DEF=Homo sapiens cDNA: FLJ21376.1s, clone COL0323.1 /FEA=mRNA /DB_XREF=gi:10437464 /UG=Hs.308748 Homo sapiens cDNA AK025029 1.37
 HGC6.2 protein NM_014356 1.37
 immunoglobulin heavy constant mu AF015124 1.37
 KIAA0241 protein A1797458 1.37
 GATA binding protein 3 A1796169 1.37

serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsin, antitrypsin), member 10	NM_016186	1.37
5-hydroxytryptamine (serotonin) receptor 3B	NM_006028	1.37
gb:U1166.1 / DEF=Human SH3 domain-containing protein SH3P17 / DB_XREF=gi:14; U8166	U8166	1.37
Consensus includes gb:AK024784.1 / DEF=Homo sapiens cDNA: FLJ21131 fls, done CAS06355, /FEA=mrna /DB_XREF=gi:10437168 /JUG=Hs.28264 Homo sapiens mRI AK024784	NM_014380	1.37
nerve growth factor receptor (TNFRSF16) associated protein 1	NM_002004	1.37
1 factor (complement)	NM_021908	1.37
suppression of tumorigenicity 7	NM_024995	1.37
hypothetical protein FLJ12616	AA209332	1.37
optic atrophy 1 (autosomal dominant)	A089655	1.37
hypothetical protein DKFZp547M136 similar to widely-interspaced zinc finger motifs	NM_017434	1.37
dual oxidase 1	NM_004984	1.37
kinase family member 5A	NM_014419	1.37
soggy-1 gene	AA052084	1.37
KIAA1001 protein	AA582460	1.37
ribosomal protein L5	BC001777	1.37
hippocalcin	A005317	1.37
TAF _{II} -like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65 kD	NM_021732	1.37
hypothetical protein PP5395	AL17663	1.37
transcription factor 3 (E2A Immunoglobulin enhancer binding factors E12/E47)	NM_004519	1.37
potassium voltage-gated channel, KQT-like subfamily, member 3	X81637	1.37
Consensus includes gb:X81637.1 / DEF=H sapiens clathrin light chain b gene, /FEA=mrna /DB_XREF=gi:963046 /JUG=Hs.73919 clathrin, light polypeptide (Lcb)	AL080178	1.37
myoblastin related protein 8	NM_014158	1.37
HSPC087 protein	NM_022367	1.37
putative metalloproteinase (family M19)	BF337329	1.37
NGF-A binding protein 2 (EGR1 binding protein 2)	Z22969	1.37
CD163 antigen	NM_001145	1.37
angiotensin, ribonuclease, RNase A family, 5	BG545288	1.37
fibronectin, B beta polypeptide	NM_017848	1.37
hypothetical protein FLJ20736	S69899	1.37
dopamine receptor D2	AA005141	1.37
met proto-oncogene (hepatocyte growth factor receptor)	NM_016307	1.37
paired related homeobox protein	BC002828	1.37
gb:BC002828.1 / DEF=Homo sapiens, Similar to fetal hypothetical protein	NM_002895	1.37
small inducible cytokine subfamily C, member 1 (lymphotactin)	U27329	1.37
foscytiferase 5 (alpha (1.3) foscytiferase)	NM_005558	1.37
lactidlin 1	A979087	1.37
transport-secretion protein 2.2,	BG236351	1.37
Consensus includes gb:BG236351 /FEA=EST /DB_XREF=est:na28112.1 /CLONE=IMAGE.4142350 /JUG=Hs.148161 ESTs	NM_015343	1.37
hypothetical protein	BF032808	1.37
Consensus includes gb:BF032808 /FEA=EST /DB_XREF=gi:10740520 /DB_XREF=est:801455629F1 /CLONE=IMAGE.3859303 /JUG=Hs.266784 ESTs	NM_006344	1.37
macrophage lectin 2 (calcium dependent)	NM_006344	1.37
Consensus includes gb:AK000834.1 / DEF=Homo sapiens cDNA FLJ20827 fls, clone ADKA03543, /FEA=mrna /DB_XREF=gi:7021152 /JUG=Hs.306409 Homo sapiens cD AK000834	NM_014898	1.37
KIAA0981 protein	NM_020215	1.37
hypothetical protein DKFZp761F2014	BF003134	1.37
chloride channel, calcium activated, family member 2	NM_001585	1.37
chromosome 22 open reading frame 1	AF006982	1.37
Consensus includes gb:AK002179.1 / DEF=Homo sapiens cDNA FLJ11317 fls, clone PLACE1010261, moderately similar to SEGREGATION DISTORTER PROTEIN, /FEA AK002179	NM_017851	1.37
hypothetical protein FLJ20509	NM_012407	1.37
protein kinase C, alpha binding protein	NM_003458	1.37
gb:NM_003458.2 / DEF=Homo sapiens high-mobility group (nonhistone chromosomal) protein isoform I-C (HMGIC), mRNA, /FEA=mrna /GEN=HMGIC /PROD=high-mobility group (nonhistone chromosomal) protein isoform I-C (HMGIC), mRNA, /FEA=mrna /DB_XREF=gi:10437464 /JUG=Hs.306748 Homo sapiens cD AK025029	NM_002188	1.37
DKFZP564A122 protein	AA521269	1.37
desmocollin 3	A1797281	1.37
VEGF nerve growth factor inducible	NM_003378	1.37
peptidyl arginine deiminase, type V	AF228067	1.37
tissue-specific heterogeneous nuclear ribonucleoprotein G-T	AF069882	1.37
paired box gene 5 (B-cell lineage specific activator protein)	BF510892	1.37
basoon (presynaptic cytomatrix protein)	NM_003458	1.37
interleukin 13	NM_002188	1.37
serine/threonine kinase 29	AJ008701	1.37
Consensus includes gb:AK025029.1 / DEF=Homo sapiens cDNA: FLJ21376 fls, clone COL03231, /FEA=mrna /DB_XREF=gi:10437464 /JUG=Hs.306748 Homo sapiens cD AK025029	AW007335	1.37
voltage-gated sodium channel beta-3 subunit (scn3b gene)	NM_025089	1.37
hypothetical protein FLJ23497	NM_016584	1.37
BM88 antigen		1.37

calmodulin-like 3
KIAA0375 gene product
Consensus includes gb:AL080072.1 /DEF=Homo sapiens mRNA; cDNA DKF Zp564M0616 (from clone DKF Zp564M0616) /FEA=mRNA /DB_XREF=gi:3282482 /UG=Hs.21 AL080072
ectonucleoside triphosphate diphosphohydrolase 5
hypothetical protein MGC2477
S100 calcium binding protein A9 (calgranulin B)
KIAA1077 protein
runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)
hypothetical protein PRO0800
solute carrier family 16 (monocarboxylic acid transporters), member 5
Consensus includes gb:Z21987.1 /DEF=H sapiens partial cDNA for homologue of mPOU homeobox protein. /FEA=mRNA /PROD=mPOU homeobox protein /DB_XREF=gi:Z21987
vavlin 3
rearranged L-myc fusion sequence
2'-5'-oligoadenylate synthetase-like
KIAA0940 protein
potential tumor suppressor
plasma glutamate carboxypeptidase
hypothetical protein FLJ22714
angiotensin 1
leucine-rich and death domain containing
small inducible cytokine A5 (RANTES)
hypothetical protein FLJ13511
jagged 1 (Alagille syndrome)
Rho GTPase activating protein 5
matrix metalloproteinase 16 (membrane-inserted)
gb:NM_007116.1 /DEF=Homo sapiens tenascin XA (TNXA). mRNA. /FEA=mRNA /GEN=TNXA /PROD=tenascin XA /DB_XREF=gi:6005907 /UG=Hs.283750 tenascin XA // NM_007116
BH-prolactin (brain-heart)
paired basic amino acid cleaving system 4
RNA binding motif, single stranded interacting protein 3
Consensus includes gb:BE551219 /FEA=EST /DB_XREF=gi:9792911 /DB_XREF=est:7656011.x1 /CLONE=IMAGE:323221 /UG=Hs.16762 Homo sapiens mRNA; cDNA I BE551219
group XII secreted phospholipase A2
matrix metalloproteinase-like 1
nuclear receptor subfamily 2, group E, member 3
iriparite motif-containing 10
DKF ZP5660193 protein
cyclin-dependent kinase inhibitor 2C (p18, inhibits CDK4)
phosphodiesterase 11A
gonadotropin-regulated testicular RNA helicase
cyclin-dependent kinase-like 2 (CDC2-related kinase)
lucosyltransferase 6 (alpha (1,3) glucosyltransferase)
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3
transcription factor 3 (E2A immunoglobulin enhancer binding factor E12/E47)
KIAA1117 protein
protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform
alcohol dehydrogenase 1B (class I), beta polypeptide
hypothetical protein FLJ11136
EGF-like domain, multiple 5
chromosome 1 open reading frame 9
Kelch motif containing protein
hypothetical protein
Consensus includes gb:AV691491 /FEA=EST /DB_XREF=gi:10293354 /DB_XREF=est:AV691491 /CLONE=GKCEEB10 /UG=Hs.85335 Homo sapiens mRNA; cDNA DKF ZP5691491
ephin-B2
E2F transcription factor 3
prodynorphin
hypothetical protein FLJ13241
Consensus includes gb:BG389073 /FEA=EST /DB_XREF=gi:13282519 /DB_XREF=est:60241488F1 /CLONE=IMAGE:4523201 /UG=Hs.151010 ESTs
ets variant gene 1
Notch homolog 3 (Drosophila)
cold autoimmune syndrome 1
anterior gradient 2 (Xenopus laevis) homolog
paraoxonase 2
heat shock 70kD protein 1A
tumor necrosis factor (ligand) superfamily, member 10

M58026 1.37
AB002373 1.37
NM_001249 1.37
BC001434 1.37
NM_002965 1.37
BE500877 1.37
L34596 1.37
NM_016592 1.37
AA705628 1.37
Z21987 1.37
NM_018399 1.37
NM_012421 1.37
NM_003733 1.37
NM_014912 1.37
NM_013437 1.37
NM_008102 1.37
NM_025050 1.37
U63508 1.37
NM_016494 1.37
NM_002985 1.37
AK023573 1.37
U61276 1.37
AY044631 1.37
NM_022564 1.37
NM_007116 1.37
NM_002589 1.37
D87894 1.37
NM_014483 1.37
BE551219 1.36
NM_030821 1.36
NM_004142 1.36
NM_016346 1.36
AF220123 1.36
BC005398 1.36
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NM_016953 1.36
NM_013284 1.36
NM_003948 1.36
U27336 1.36
AF009644 1.36
AB71234 1.36
AL162056 1.36
AF070670 1.36
AF153821 1.36
A497781 1.36
BF110421 1.36
AC004770 1.36
NM_014458 1.36
NM_019079 1.36
U16797 1.36
AB03363 1.36
NM_024411 1.36
NM_025088 1.36
BG389073 1.36
AC004857 1.36
NM_000435 1.36
AK027194 1.36
AF088867 1.36
AF001602 1.36
NM_005345 1.36
NM_003810 1.36

fizzled-related protein	U91903	1.36
G protein-coupled receptor 86	NM_023914	1.36
protein disulfide isomerase-related protein	BE910010	1.36
hypothetical protein FLJ23058	NM_024696	1.36
solute carrier family 9 (sodium/hydrogen exchanger), isoform 2	NM_003048	1.36
homolog of yeast ATP12	AW118808	1.36
cytochrome b-5	M22865	1.36
LCU7-like (S. cerevisiae)	NM_018032	1.36
scrapie responsive protein 1	NM_007281	1.36
discolpin domain receptor family, member 1	NM_001954	1.36
Bedilius subtilis /REF=L38424 /DEF=B subtilis dapB, joiF, joiG genes corresponding to nucleotides 2055-2578 of L38424 /LEN=1931 (-5, -M, -3 represent transcript regions)	L38424	1.36
hypothetical protein	A1746515	1.36
relaxin 1 (H1)	BC005956	1.36
KIAA0103 gene product	NM_014673	1.36
TAF6-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65 kD	AI005317	1.36
prolactin receptor	AF166329	1.36
G antigen 3	NM_001473	1.36
Ca2-B-M (murine) ectropic retroviral transforming sequence b	NM_004351	1.36
vacuolar protein sorting 33B (yeast)	AI672363	1.36
cyclic nucleotide gated channel alpha 1	NM_000087	1.36
SMT3 suppressor of nilf two 3 homolog 2 (yeast)	AV726646	1.36
caveolin 1, caveolae protein, 22kD	NM_001753	1.36
paraoxonase 2	NM_000305	1.36
fibroblast growth factor 13	NM_004114	1.36
oxytocin receptor	NM_000916	1.36
peroxisome proliferative activated receptor, gamma	NM_015869	1.36
Consensus includes gb:AF052090.1 /DEF=Homo sapiens clone 23950 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360396 /UG=Hs.106620 Homo sapiens clone 23950	AF052090	1.36
arylsulfatase B	NM_000046	1.36
slaytransferase 4A (beta-galactosidase alpha-2,3-sialyltransferase)	NM_003033	1.36
protein phosphatase 1, regulatory (inhibitor) subunit 3A (glycogen and sarcoplasmic reticulum binding subunit, skeletal muscle)	NM_002711	1.36
aminohexopate-semialdehyde dehydrogenase-phosphopantetheinyl transferase	AF302110	1.36
G antigen 4	NM_001474	1.36
Consensus includes gb:AL049233.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564A023 (from clone DKFZp564A023). /FEA=mRNA /DB_XREF=gi:4499987 /UG=Hs.4224	AL049233	1.36
aldo-keto reductase family 1, member D1 (delta 4-3-ketosteroid-5-beta-reductase)	NM_005689	1.36
peroxisome proliferative activated receptor, delta	NM_006238	1.36
coagulation factor XIII, A1 polypeptide	NM_000129	1.36
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 1	U73396	1.36
Consensus includes gb:BC002892.1 /DEF=Homo sapiens, clone IMAGE:3940152, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3940152) /DB_XI	BC002892	1.36
likely ortholog of mouse ganglioside-induced differentiation-associated-protein 2	NM_017686	1.36
CD8 antigen, alpha polypeptide (p32)	AW006735	1.36
chromosome 1 open reading frame 15	AF288395	1.36
cell division cycle 2-like 5 (cholinesterase-related cell division controller)	AJ297710	1.36
hypothetical protein MGC5601	NM_025247	1.36
hypothetical protein FLJ22251	NM_025158	1.36
chromosome 5 open reading frame 3	NM_016691	1.36
Kruppel-like factor 4 (gut)	BF514079	1.36
Consensus includes gb:U80771.1 /DEF=Human EST clone 25267 mariner transposon Hsmar1 sequence. /FEA=mRNA /DB_XREF=gi:2231374 /UG=Hs.245410 Human ES	U80771	1.36
phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 duncle homolog, Drosophila)	AF012073	1.36
hepatitis delta antigen-interacting protein A	A583181	1.36
tripartite motif-containing 2	BC005016	1.36
likely ortholog of mouse variant polyadenylation protein CSTF-64	A672408	1.36
hypothetical protein FLJ20602	NM_017896	1.36
Consensus includes gb:AK022892.1 /DEF=Homo sapiens cDNA FLJ12830 fls, clone NT2RP2003073. /FEA=mRNA /DB_XREF=gi:10434549 /UG=Hs.123158 Homo sapi	AK022892	1.36
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	D42072	1.36
hypothetical protein FLJ12691	NM_024572	1.36
ATPase, Na+/K+ transporting, alpha 2 (+) polypeptide	AW440492	1.36
Consensus includes gb:AJ243936.1 /DEF=Homo sapiens mRNA for G16 protein located in the class III region of the major histocompatibility complex. /FEA=m	AJ243936	1.36
Consensus includes gb:AL049325.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564D036 (from clone DKFZp564D036). /FEA=mRNA /DB_XREF=gi:4500098 /UG=Hs.1124	AL049325	1.36
zinc finger protein 44 (KIX 7)	AI758888	1.36
N amino acid transporter 3	NM_018018	1.36
gap junction protein, alpha 4, 37kD (connexin 37)	M86789	1.36
small inducible cytokine subfamily A (Cys-Cys), member 22	NM_002990	1.36
melatonin receptor 1B	NM_005859	1.36

siaphorin (gpl.115, leukostellin, CD43)	X52075	1.36
spermidine/spermine N1-acetyltransferase	NM_002870	1.36
Consensus includes gb:AL031714 /DEF=Human DNA sequence from clone LA16-35887 on chromosome 16 Contains the UBE21 gene for ubiquitin-conjugating enzyme E1	AL031714	1.36
doublecortin; lissencephaly, X-linked (doublecortin)	NM_000555	1.36
nuclear factor (erythroid-derived 2), 45kD	L13974	1.36
RNA binding motif protein 6	NM_005777	1.36
supervillin	NM_003174	1.36
KIAA0317 gene product	NM_014821	1.36
helicase-mot	NM_030621	1.36
rabaptin-5	AF098638	1.36
hypothetical protein FLJ13055	AA209463	1.36
Consensus includes gb:AK025323.1 /DEF=Homo sapiens cDNA: FLJ21670 fls, clone COL09010, /FEA=mRNA /DB_XREF=gi:10437816 /UG=Hs.230999 Homo sapiens cDNA: AK025323	N32185	1.36
Cluster incl. N32185/Y2510.11 Homo sapiens cDNA, 3 end /clone=IMAGE:272275 /clone_end=3 /gb=N32185 /gi=1152584 /ug=Hs.120769 /len=600	AF320053	1.36
v-myc myelocytomatosis viral related oncogene, neuroblastoma derived (avian)	NM_017892	1.36
formin-binding protein 3	NM_014055	1.36
CDV-1 protein	AW298219	1.36
KIAA1016 protein	NM_017417	1.36
UDP-N-acetyl-alpha-D-galactosamine polypeptide N-acetyl-galactosaminyltransferase 8 (GalNAc-T8)	NM_006141	1.36
dynein, cytoplasmic, light intermediate polypeptide 2	AK022063	1.36
Consensus includes gb:AK022063.1 /DEF=Homo sapiens cDNA FLJ12001 fls, clone HEMBB1001535, /FEA=mRNA /DB_XREF=gi:10433383 /UG=Hs.278412 Homo sapiens cDNA: FL	BG289001	1.36
Consensus includes gb:BG289001 /FEA=EST /DB_XREF=gi:13044404 /DB_XREF=est:602381282F1 /CLONE=IMAGE:4499078 /UG=Hs.101282 Homo sapiens cDNA: FL	AL575403	1.36
KIAA0820 protein	NM_005433	1.36
v-yes-1 Yamaguchi sarcoma viral oncogene homolog 1	NM_002309	1.36
leukemia inhibitory factor (proliferation differentiation factor)	NM_025139	1.36
Consensus includes gb:AF070625.1 /DEF=Homo sapiens clone 24734 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3283891 /UG=Hs.12440 Homo sapiens clone 24734, AF070625	BC002599	1.36
hypothetical protein FLJ12584	NM_012118	1.36
coricotropin releasing hormone	BC005303	1.36
CCRA carbon catabolite repression 4-like (S. cerevisiae)	AW301841	1.36
prolamine 2	NM_014565	1.36
RAB2, member RAS oncogene family	X68465	1.36
olfactory receptor, family 1, subfamily A, member 1	BF593908	1.36
adenosine A1 receptor	NM_003156	1.36
TATA element modulatory factor 1	BC004134	1.36
stromal interaction molecule 1	AK024606	1.36
advinin	BF229395	1.36
Consensus includes gb:AK024606.1 /DEF=Homo sapiens cDNA: FLJ20953 fls, clone ADSE01978, /FEA=mRNA /DB_XREF=gi:10436920 /UG=Hs.306697 Homo sapiens cDNA: AK024606	BF229395	1.36
Consensus includes gb:BF229395 /FEA=EST /DB_XREF=gi:11166291 /DB_XREF=est:MR1-C10019-260900-001-a11 /UG=Hs.159454 ESTs	X17247	1.36
Consensus includes gb:AW265295 /FEA=EST /DB_XREF=gi:6701931 /DB_XREF=est:UH-H-B12-ah1-1-2-0-U1.s1 /CLONE=IMAGE:2728079 /UG=Hs.212522 ESTs, Highly	NM_004770	1.36
sialyltransferase 1 (beta-galactoside alpha-2,6-sialyltransferase)	NM_008485	1.36
potassium voltage-gated channel, Shab-related subfamily, member 2	NM_022562	1.36
fibulin 1	NM_015874	1.36
growth hormone 1	Z58820	1.36
H-2K binding factor-2	NM_024631	1.36
Consensus includes gb:X87875 /DEF=H.sapiens EP4 prostaglandin receptor pseudogene c16g7 /FEA=CDS /DB_XREF=gi:1359737 /UG=Hs.247972 H.sapiens EP4 prost	M74777	1.36
Interleukin 17 (cytotoxic T-lymphocyte-associated serine esterase 8)	M23254	1.36
hypothetical protein FLJ23342	L32867	1.36
dipeptidylpeptidase IV (CD26, adenosine deaminase complexing protein 2)	NM_025099	1.36
calpain 2, (mII) large subunit	BC026366	1.36
sialyltransferase 8 (alpha-N-acetylneuraminic acid: alpha-2,8-sialyltransferase, GD3 synthase) A	NM_016584	1.36
hypothetical protein FLJ22170	NM_022353	1.36
KH-type splicing regulatory protein (FUSE binding protein 2)	NM_003426	1.36
interleukin 23, alpha subunit p19	NM_013270	1.36
putative sialylglycoprotease type 2	AL162081	1.36
zinc finger protein 74 (Czsf52)	NM_001987	1.36
testes-specific protease 50	NM_000420	1.36
RAB14, member RAS oncogene family	AK024553	1.36
eta variant gene 6 (TEL oncogene)	NM_023908	1.36
Kell blood group	U17105	1.36
Consensus includes gb:AK024553.1 /DEF=Homo sapiens cDNA: FLJ20900 fls, clone ADKA03593, /FEA=mRNA /DB_XREF=gi:10436861 /UG=Hs.306687 Homo sapiens cDNA: AK024553	A1382018	1.36
v-ret reticulendotheliosis viral oncogene homolog (avian)		
cyclin F		
cylindromatosis (urban tumor syndrome)		
carbamite palmitoyltransferase 1, liver		

KIAA0073 protein	AK025679	1.35
DKFZP564A122 protein	AL110243	1.35
gsk3j phosphoprotein 2	NM_016548	1.35
platelet factor 4	NM_002619	1.35
retinoic acid receptor, beta	NM_016152	1.35
limbic system-associated membrane protein	NM_002338	1.35
Hermansky-Pudlak syndrome	NM_000195	1.35
calcium-sensing receptor (hypocalcemic hyperparathyroidism)	D50855	1.35
hypothetical protein FLJ20202	NM_017709	1.35
adenylate cyclase 9	NM_001116	1.35
meningioma expressed antigen 5 (hyaluronidase)	NM_012215	1.35
discs, large (Drosophila) homolog 4	NM_001365	1.35
bromodomain adjacent to zinc finger domain, 2A	BE513659	1.35
sialic acid binding Ig-like lectin 7	AJ130713	1.35
DKFZP566O1646 protein	AF25793	1.35
neurochondrin	AB018740	1.35
rebaptin-5	NM_004703	1.35
hypothetical protein MCC5347	BE680973	1.35
Consensus includes gb:AV720842 /FEA=EST /DB_XREF=gi:10817994 /DB_XREF=est:AV720842 /CLONE=GLCCOH05 /UG=Hs.282435 ESTs	AV720842	1.35
protein kinase, X-linked	NM_005044	1.35
neurexin 1	AB035356	1.35
solute carrier family 2 (facilitated glucose transporter), member 3	AL110298	1.35
chromodomain helicase DNA binding protein 3	NM_001272	1.35
myosin binding protein H	NM_004997	1.35
trihydrocidic repeat containing 9	AK025084	1.35
lipase, hormone-sensitive	NM_005357	1.35
bradykinin receptor B2	NM_000823	1.35
laminin, gamma 1 (formerly LAMB2)	NM_002283	1.35
TBP-interacting protein	AB014567	1.35
tumor necrosis factor (ligand) superfamily, member 11	AF053712	1.35
HOA174	NM_016484	1.35
glycine receptor, alpha 3	NM_006529	1.35
hypothetical protein DKFZP547M136 similar to widely-interspaced zinc finger motifs	W67467	1.35
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 7 (18kD, B18)	M33374	1.35
hypothetical protein FLJ20489	AW148696	1.35
cell division cycle 42 (GTP binding protein, 25kD)	BC003682	1.35
lucosyltransferase 4 (alpha 1.3) fucosyltransferase, myeloid-specific)	AF305083	1.35
peripheral benzodiazepine receptor-associated protein 1	NM_004756	1.35
hypothetical protein FLJ20087	NM_017662	1.35
Consensus includes gb:AV712602 /FEA=EST /DB_XREF=gi:10731908 /DB_XREF=est:AV712602 /CLONE=DCAACD04 /UG=Hs.5957 Homo sapiens clone 24416 mRNA	AV712602	1.35
purinergic receptor P2X-like 1, orphan receptor	AF065385	1.35
ATP-binding cassette, sub-family A (ABC1), member 4	U88667	1.35
regulator of G-protein signalling 1	NM_002922	1.35
muscle disease-related protein	NM_016298	1.35
arylsulfatase desacylase (esterase)	NM_001086	1.35
epithelial V-like antigen 1	NM_005797	1.35
LIM homeobox transcription factor 1, beta	NM_002316	1.35
v-akt murine thymoma viral oncogene homolog 2	NM_001038	1.35
SRY (sex determining region Y)-box 4	M77198	1.35
hypothetical protein PR02325	BG528420	1.35
Consensus includes gb:BG540048 /FEA=EST /DB_XREF=gi:13532281 /DB_XREF=est:602568675F1 /CLONE=IMAGE:4693242 /UG=Hs.293982 ESTs	NM_018523	1.35
xylokinase (H. influenzae) homolog	BG540048	1.35
Consensus includes gb:AL567940 /FEA=EST /DB_XREF=gi:12921802 /DB_XREF=est:AL567940 /CLONE=CS0DF036YK19 (3 prime) /UG=Hs.7967 ESTs	NM_005108	1.35
caspase recruitment domain protein 14	NM_024110	1.35
pulsative ankyrin-repeat containing protein	NM_015623	1.35
CD84 antigen (leukocyte antigen)	AF054818	1.35
crystallin, beta A4	NM_001886	1.35
cholinergic receptor, muscarinic 5	NM_012125	1.35
Consensus includes gb:AF037219.1 /DEF=Homo sapiens PIX1 mRNA sequence, /FEA=mRNA /DB_XREF=gi:4090971 /UG=Hs.301667 Homo sapiens mRNA; cDNA DKF1	AF037219	1.35
FXYD domain-containing ion transport regulator 2	X86400	1.35
KIAA1042 protein	AK000754	1.35
BAL1-associated protein 2	BC002495	1.35

mitogen-activated protein kinase 8	U34822	1.35
carboxypeptidase, vitellogenic-like	NM_031311	1.35
hypothetical protein PRO2730	N82524	1.35
ADP-ribosyltransferase (NAD ⁺ ; poly (ADP-ribose) polymerase)-like 1	NM_006437	1.35
tenascin XB	M25813	1.35
kinasin family member 58	BF223224	1.35
tumor necrosis factor, alpha-induced protein 2	NM_006291	1.35
calcium channel, voltage-dependent, gamma subunit 5	NM_014404	1.35
adrenergic, alpha-1D-, receptor	M76448	1.35
proedlin P factor, complement	NM_002621	1.35
neutrophil cytosolic factor 2 (65kD, chronic granulomatous disease, autosomal 2)	BC001606	1.35
KIA0756 protein	AB217777	1.35
SCAN domain-containing 2	AF244812	1.35
Consensus includes gb:A185160 /FEA=EST /DB_XREF=gi:3735798 /DB_XREF=est:q30g10.s1 /CLONE=IMAGE:1740546 /UG=Hs.326248 Homo sapiens cDNA: FLJ220	AF185160	1.35
synovial sarcoma, X breakpoint 3	S82471	1.35
zinc finger protein 291	AF118814	1.35
hypothetical protein FLJ20581	NM_017868	1.35
KIAA0508 protein	AB007975	1.35
echinoderm microtubule-associated protein-like	NM_004434	1.35
paired box gene 5 (B-cell lineage specific activator protein)	NM_016734	1.35
mitogen-activated protein kinase kinase 7	AF006689	1.35
thyroglobulin	NM_003235	1.35
TAL1 (SCL) interrupting locus	NM_003035	1.35
chromosome 12 open reading frame 6	NM_020367	1.35
adrenergic, beta, receptor kinase 1	NM_001819	1.35
uroplakin 3	NM_006953	1.35
corticotropin releasing hormone receptor 1	AF180301	1.35
HIV-1 Tat interactive protein 2, 30 kDa	AF092085	1.35
cancer testis antigen	AJ275978	1.35
glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)	XG9936	1.35
hypothetical protein FLJ12681	AL515916	1.35
Consensus includes gb:AL551048 /FEA=EST /DB_XREF=gi:12888616 /DB_XREF=est:AL551048 /CLONE=CS00D066SYN1.3 (5 prime) /UG=Hs.332742 Human DNA from cl	AL551046	1.35
gb:AF165312.1 /DEF=Homo sapiens pre-T-cell receptor alpha chain (PTA) mRNA, alternatively spliced, complete cds. /FEA=CD5 /GEN=PTA /PROD=pre-T-cell receptor a	AF165312	1.35
hypothetical protein FLJ21195 similar to protein related to DAC and cerberus	NM_022469	1.35
hypothetical protein FLJ21324	AW449022	1.35
potassium inwardly-rectifying channel subfamily J, member 13	AJ007557	1.35
Consensus includes gb:AF005081.1 /DEF=Homo sapiens skin-specific protein (xp32) mRNA, partial cds. /FEA=mRNA /GEN=xp32 /PROD=skin-specific protein /DB_XREF=	NM_005764	1.35
epithelial protein up-regulated in carcinoma, membrane associated protein 17	M24915	1.35
CD44 antigen (homolog function and Indian blood group system)	NM_024812	1.35
hypothetical protein FLJ12015	U07969	1.35
cadherin 17, LI cadherin (liver-intestine)	NM_007327	1.35
glutamate receptor, ionotropic, N-methyl D-aspartate 1	NM_002784	1.35
RuvB-like 1 (E. coli)	NM_005393	1.35
plexin B3	A056359	1.35
microtubule-associated protein tau	NM_000266	1.35
Norrie disease (pseudoglioma)	AL161996	1.35
ATPase, Class VI, type 11A	NM_030971	1.35
similar to rat tricarboxylate carrier-like protein	N85363	1.35
cyclin-dependent kinase inhibitor 1C (p57, Kip2)	NM_016529	1.35
ATPase, aminophospholipid transporter-like, Class 1, type BA, member 2	NM_016236	1.35
alanine-glyoxylate aminotransferase (oxalosis I; hyperoxaluria I; glycolicaciduria; serine-pyruvate aminotransferase)	NM_005371	1.35
methyltransferase-like 1	AL554008	1.35
G protein-coupled receptor 58	NM_001174	1.35
Rho GTPase activating protein 6	NM_001450	1.35
four and a half LIM domains 2	NM_017918	1.35
hypothetical protein FLJ20847	NM_014847	1.35
KIAA0144 gene product	NM_002353	1.35
tumor-associated calcium signal transducer 2	AF038181	1.35
Consensus includes gb:AF038181.1 /DEF=Homo sapiens clone 23607 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2795901 /UG=Hs.167825 Homo sapiens clone 23607	AF038181	1.35
triple functional domain (PTPRF interacting)	AA716657	1.35
presenilin 2 (Alzheimer disease 4)	NM_000111	1.35
solite carrier family 28, member 3	NM_015839	1.35
ficolin (collagen/fibrinogen domain-containing lectin) 2 (hucolin)		1.35

centaurin, gamma 2	NM_014914	1.35
hypothetical protein	NM_013310	1.35
a disintegrin-like and metalloprotease (repolysin type) with thrombospondin type 1 motif, 8	NM_007037	1.35
putative opioid receptor, neuromedin K (neurokinin B) receptor-like	NM_006879	1.35
calcium channel, voltage-dependent, L type, alpha 1B subunit	NM_000718	1.35
RAR-related orphan receptor B	NM_006914	1.35
ankyrin repeat domain 3	NM_020639	1.35
cytochrome P450, subfamily IVF, polypeptide 2	D26460	1.35
Consensus includes gb:AA551075 /FEA=EST /DB_XREF=est:nk74h06.s1 /CLONE=IMAGE:1019291 /UG=Hs.109438 Homo sapiens clone 24775 r	AA551075	1.35
hypothetical protein FLJ20034	NM_017630	1.35
myotubularin related protein 1	AF057354	1.35
SPR1 protein	NM_014069	1.35
dual-specific tyrosine-(Y)-phosphorylation regulated kinase 1B	NM_004714	1.35
actinin, alpha 2	NM_001103	1.35
KIAA1035 protein	AB028958	1.35
chemokine binding protein 2	NM_001296	1.35
hypothetical protein MGC2592	NM_121891	1.35
Consensus includes gb:A050036 /FEA=EST /DB_XREF=est:an31a09.x1 /CLONE=IMAGE:1700248 /UG=Hs.190616 ESTs	A050036	1.35
KIAA0419 gene product	NM_014711	1.35
functional adhesion molecule 2	NM_021219	1.35
zinc finger protein 93 homolog (mouse)	BC002863	1.35
bromodomain-containing 2	S78711	1.35
resin (Reed-Steinberg cell-expressed intermediate filament-associated protein)	M97501	1.35
Gardner-Rasheed feline sarcoma viral (v-fgr) oncogene homolog	BC002836	1.35
endothelial differentiation, sphingolipid G-protein-coupled receptor, 1	NM_001400	1.35
neurobeachin	NM_013678	1.35
cardiac ankyrin repeat protein	NM_014391	1.35
Consensus includes gb:AW975638 /FEA=EST /DB_XREF=est:EST387747 /UG=Hs.293490 ESTs	AW975638	1.35
KIAA0241 protein	BE505338	1.35
hemochromatosis	AF144243	1.35
Consensus includes gb:AL122056.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434P088 (from clone DKFZp434P088) /FEA=mRNA /DB_XREF=gi:5093253 /UG=Hs.27451	AL122056	1.35
nucleoporin 88KD	Y08613	1.35
transporlin-SR	AK022910	1.35
NK-receptor	AF022048	1.35
hypothetical protein FLJ20079	NM_017856	1.35
homeo box B9	X16172	1.35
surfeit 2	NM_017503	1.35
Consensus includes gb:BE873723 /FEA=EST /DB_XREF=gi:1003426 /DB_XREF=est:7d9n10.x1 /CLONE=IMAGE:3279235 /UG=Hs.82719 Homo sapiens mRNA; cDNA BE873723	BE873723	1.35
glycophorin A (includes MN blood group)	NM_020999	1.35
AKAP-binding sperm protein ropporin	NM_017578	1.35
regulator of G-protein signalling 3	NM_021106	1.35
Yea-associated protein 1, 65 kDa	A1745185	1.35
M-phase phosphoprotein 6	NM_005782	1.35
tripartite motif-containing 22	AA083478	1.35
homeo box A11	NM_005523	1.35
glucosyltransferase 8 (alpha (1,6) glucosyltransferase)	NM_004480	1.35
claudin 4	NM_001305	1.35
G protein-coupled receptor 32	NM_001506	1.35
leucine rich repeat (in FLI) interacting protein 2	NM_017724	1.35
serine (or cysteine) proteinase inhibitor, clade C (antithrombin), member 1	D29832	1.35
chromosome 21 open reading frame 7	NM_020152	1.35
DKFZP434N161 protein	NM_015493	1.35
guanylate binding protein 2, interferon-inducible	NM_004120	1.35
gamma-aminobutyric acid (GABA) A receptor, pi	NM_014211	1.35
pim-2 oncogene	NM_006875	1.35
DKFZP586111 protein	AL050131	1.35
natural killer cell receptor, immunoglobulin superfamily member	NM_007053	1.35
insulin-like growth factor binding protein 4	NM_001552	1.35
Consensus includes gb:A26304 /FEA=EST /DB_XREF=gi:3871247 /DB_XREF=est:az29a03.x1 /CLONE=IMAGE:2028316 /UG=Hs.13438 Homo sapiens clone 24626 mf	A26304	1.35
hypothetical protein	NM_019062	1.35
myelopoietin	NM_005775	1.35
Consensus includes gb:AK025352.1 /DEF=Homo sapiens cDNA: FLJ21699 fs. clone COL09829 /FEA=mRNA /DB_XREF=gi:10437851 /UG=Hs.306804 Homo sapiens cDNA: AK025352	AK025352	1.35

diacylglycerol kinase, alpha (80kD)	NM_001345	1.35
CD164 antigen, siomucin	NM_006016	1.35
KIAA0773 gene product	NM_014690	1.35
organic cationic transporter-like 3	NM_004256	1.35
Consensus includes gb:AV699347 /FEA=EST /DB_XREF=est:AV699347 /CLONE=GKCBPH07 /UG=Hs.83623 nuclear receptor subfamily 1, group M/4690	M04690	1.35
pregnancy specific beta-1-glycoprotein 9	AL583811	1.35
Wilms' tumour 1-associated protein	AF000426	1.35
lymphocyte antigen 117	NM_024803	1.34
hypothetical protein FLJ12665	NM_024803	1.34
Consensus includes gb:AU145746 /FEA=EST /DB_XREF=gi:11007267 /DB_XREF=est:AU145746 /CLONE=HEMBA1005679 /UG=Hs.82193 esterase D/ormylglutathione t	AU145746	1.34
Consensus includes gb:AL359032.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 1968422. /FEA=mRNA /PROD=ITGBL1, integrin beta-like 1 /D	AL359032	1.34
nuclear receptor subfamily 2, group F, member 1	BC004154	1.34
cyclic nucleotide gated channel alpha 3	NM_001298	1.34
SRY (sex determining region Y)-box 2	AL56682	1.34
phosphoryl-CoA hydroxylase interacting protein	NM_014759	1.34
opioid receptor, delta 1	NM_000911	1.34
hypothetical protein DKFZp434N074	AL096732	1.34
sialic acid binding Ig-like lectin 5	NM_003830	1.34
cysteine-rich hydrophobic domain 2	NM_012110	1.34
interleukin 9 receptor	284723	1.34
discs, large (Drosophila) homolog 5	BC002915	1.34
LBP protein; likely ortholog of mouse CRTR-1	NM_014553	1.34
cytochrome b5 reductase b5R.2	NM_016229	1.34
synapsin III	NM_003490	1.34
protein predicted by clone 23627	NM_013299	1.34
DKFZP434N093 protein	AL080144	1.34
hypothetical protein FLJ10134	NM_018004	1.34
hypothetical protein FLJ20619	NM_017904	1.34
Consensus includes gb:AL079294.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 362780. /FEA=mRNA /DB_XREF=gi:5102751 /UG=Hs.28952	AL079294	1.34
insulin-like 5	NM_005478	1.34
regulatory factor X, 2 (influences HLA class II expression)	NZ5562	1.34
Consensus includes gb:A344415 /FEA=EST /DB_XREF=gi:4081621 /DB_XREF=est:cp13b08.x1 /CLONE=IMAGE:1917879 /UG=Hs.156082 ESTs	NM_000635	1.34
stress-associated endoplasmic reticulum protein 1, ribosome associated membrane protein 4	A344415	1.34
NAG-5 protein	BG107676	1.34
dopachrome tautomerase (dopachrome delta-isomerase, tyrosine-related protein 2)	NM_016446	1.34
chemokine (C-C motif) receptor 5	NM_001922	1.34
Consensus includes gb:AU157915 /FEA=EST /DB_XREF=gi:11019436 /DB_XREF=est:AU157915 /CLONE=PLACE1010102 /UG=Hs.5518 Homo sapiens cDNA FLJ11311	NM_000579	1.34
isocitrate dehydrogenase 2 (NADP+), mitochondrial	AU157915	1.34
paired box gene 8	U52144	1.34
tripartite motif-containing 34	AI948472	1.34
hypothetical protein MGC2306	NM_021616	1.34
Consensus includes gb:BE674468 /FEA=EST /DB_XREF=gi:10035007 /DB_XREF=est:7e04e01.x1 /CLONE=IMAGE:3281496 /UG=Hs.12514 Homo sapiens clone 23688 r	BC002557	1.34
hypothetical protein PRO2037	NM_018616	1.34
Consensus includes gb:AU144530 /FEA=EST /DB_XREF=gi:11006051 /DB_XREF=est:AU144530 /CLONE=HEMBA1002189 /UG=Hs.296654 Homo sapiens cDNA FLJ11	NM_018616	1.34
KIAA1053 protein	AL117523	1.34
myeloid/lymphoid or mixed-lineage leukemia (t(11h10r)ax homolog, Drosophila); translocated to, 4	NM_005936	1.34
Consensus includes gb:AL157837 /DEF=Human DNA sequence from clone RP11-467121 on chromosome 1q31.1-31.3 Contains a ribosomal protein L23A pseudogene, EE	AL157837	1.34
catenin (cadherin-associated protein), beta 1 (88kD)	NM_001904	1.34
Consensus includes gb:W88821 /FEA=EST /DB_XREF=gi:1404293 /DB_XREF=est:zn71d12.r1 /CLONE=IMAGE:417527 /UG=Hs.288924 Homo sapiens cDNA FLJ11392	W88821	1.34
retinal short-chain dehydrogenase/reductase rslSDR3	NM_016246	1.34
RAB6B, member RAS oncogene family	AW118072	1.34
cathepsin K (pseudocystosis)	NM_000396	1.34
hypothetical protein FLJ11292	NM_018382	1.34
phosphodiesterase 8B	AK023913	1.34
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 9	NM_020249	1.34
four and a half LIM domains 1	U29538	1.34
heparan sulfate 6-O-sulfotransferase	BF512553	1.34
KIAA0143 protein	AW470003	1.34
lipidolisin	AB014531	1.34
caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)	AU719655	1.34
leptin receptor gene-related protein	AW026535	1.34
Cluster Ind. A1742455:wg43605.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2367625 /clone_end=3 /gb=A1742455 /gi=5110743 /ug=Hs.28974 /len=489	A1742455	1.34

potassium voltage-gated channel, shaker-related subfamily, beta member 3
Consensus includes gb:AV687010 /FEA=EST /DB_XREF=gi:10288873 /DB_XREF=est:AV687010 /CLONE=GKCAVC04 /UG=Hs.284651 ESTs

lipin 1
nuclear receptor subfamily 4, group A, member 2
nuclear antigen Sp100
UDP-N-acetylglucosamine-1,3-D-mannoside beta-1,4-N-acetylglucosaminyltransferase IV-homolog
hypothetical protein TCBAPO758
wingless-type MMTV integration site family, member 2B
spinocerebellar ataxia 7 (olivopontocerebellar atrophy with retinal degeneration)
Consensus includes gb:AL050378.1 /DEF=Homo sapiens mRNA, cDNA DKFZp586i1420 (from clone DKFZp586i1420), partial cds. /FEA=mRNA /GEN=DKFZp586i1420
GLI-Kruppel family member GLI2
histone acetyltransferase
opoid growth factor receptor
nuclear receptor subfamily 1, group 1, member 2
BTG family, member 3
KIAA1025 protein
retinoblastoma binding protein 4
Consensus includes gb:Z34282.1 /DEF=H sapiens (MAR11) MUC5AC mRNA for mucin (partial). /FEA=mRNA /GEN=MUC5AC /PROD=mucin /DB_XREF=gi:563384 /UG=Hs.294277 ESTs
KIAA0914 gene product
Consensus includes gb:A078169 /FEA=EST /DB_XREF=gi:3412577 /DB_XREF=est:z30d11.x1 /CLONE=IMAGE:1676853 /UG=Hs.323511 Homo sapiens cDNA: FLJ231
radixin
Consensus includes gb:AF339787.1 /DEF=Homo sapiens clone IMAGE:205688, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507325 /UG=Hs.326722 Homo sapiens cAF339787
poly (ADP-ribose) glycohydrolase
hypothetical protein FLJ23594
Consensus includes gb:BE671816 /FEA=EST /DB_XREF=gi:10032446 /DB_XREF=est:7a7d01.x1 /CLONE=IMAGE:3221857 /UG=Hs.21921 ESTs. Moderately similar to /BE671816
prefoldin 4
nuclear receptor subfamily 2, group C, member 1
Consensus includes gb:AV700891 /FEA=EST /DB_XREF=gi:10302862 /DB_XREF=est:AV700891 /CLONE=GKCBQD03 /UG=Hs.292477 ESTs
KIAA0564 protein
Consensus includes gb:AK025833.1 /DEF=Homo sapiens cDNA: FLJ22180 fls, clone HRC00936. /FEA=mRNA /DB_XREF=gi:10438467 /UG=Hs.287692 Homo sapiens cAF025833
hypothetical protein FLJ13397
calcium binding protein 1 (calbrain)
synuclein, alpha (non A4 component of amyloid precursor)
tumor protein D52-like 1
synuclein, alpha interacting protein (synphilin)
Consensus includes gb:AK022421.1 /DEF=Homo sapiens cDNA FLJ12359 fls, clone MAMMA1002355. /FEA=mRNA /DB_XREF=gi:10433814 /UG=Hs.298718 Homo sapiens cAK022421
hypothetical protein FLJ22659
hypothetical protein FLJ11029
hypocretin (orexlin) neuropeptide precursor
chromosome 12 open reading frame 5
gb:U87229.1 /DEF=Homo sapiens death adaptor molecule RAIDD-2 mRNA, complete cds. /FEA=CDS /PROD=death adaptor molecule RAIDD-2 /DB_XREF=gi:5051867 /UG=Hs.2987229
TATA box binding protein (TBP)-associated factor, RNA polymerase I, C, 110kD
Slah-interacting protein
interleukin 15
cophilin VII
fritzzed homolog 10 (Drosophila)
KIAA0856 protein
Rap1 guanine-nucleotide-exchange factor directly activated by cAMP
hypothetical protein FLJ11808
growth hormone releasing hormone
myosin light chain 2a
carbonic anhydrase I
KIAA0365 gene product
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 1
KIAA0942 protein
sphingomyelin phosphodiesterase 1, acid lysosomal (acid sphingomyelinase)
hypothetical protein FLJ20378
interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD)
mitogen-activated protein kinase 11
Consensus includes gb:AA654415 /FEA=EST /DB_XREF=gi:2590569 /DB_XREF=est:m03g09.s1 /CLONE=IMAGE:1192096 /UG=Hs.322808 ESTs. Moderately similar to A AA654415
guanylate cyclase activator 1B (relina)
protein tyrosine phosphatase, non-receptor type 12
cullin 5

likely ortholog of mouse Arfada
Consensus includes gb:A033741 /FEA=EST /DB_XREF=gi:4685071 /DB_XREF=est:128a05.x1 /CLONE=IMAGE:2242064 /UG=Hs.270364 ESTs, Moderately similar to S1
tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)
signal transducer and activator of transcription 2, 113kD
malic enzyme 3, NADP(+)-dependent, mitochondrial
ribosomal protein S6 kinase, 90kD, polypeptide 5
Consensus includes gb:AV720514 /FEA=EST /DB_XREF=gi:10817666 /DB_XREF=est:AV720514 /CLONE=GLCGS809 /UG=Hs.282721 ESTs, Weakly similar to ALU7_HI
Fc fragment of IgA receptor for
KLAA0876 protein
trypsin beta 1
ubiquitin associated protein
CCR4-NOT transcription complex, subunit 2
ceroid-lipofuscinosis, neuronal 3, juvenile (Batten, Spielmeyer-Vogt disease)
visual system homeobox 1 homolog, CHX10-like (zebrafish)
Consensus includes gb:AL110223.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434M063 (from clone DKFZp434M063) /FEA=mRNA /DB_XREF=gi:5817158 /UG=Hs.2745
integrin, alpha 5 (fibronectin receptor, alpha polypeptide)
Consensus includes gb:AL080111.1 /DEF=Homo sapiens mRNA, cDNA DKFZp586G2222 (from clone DKFZp586G2222) /FEA=mRNA /DB_XREF=gi:5262538 /UG=Hs.24
KLAA0090 protein
opsin 1 (cone pigments), medium-wave-sensitive (color blindness, deutan)
retinoic acid receptor, alpha
p21 (CDKN1A)-activated kinase 3
membrane-binding lectin (protein A) 1, pseudogene 1
hypothetical protein FLJ20276
hypothetical protein MGC2601
solute carrier family 1 (neuronal/epithelial high affinity glutamate transporter, system Xag), member 1
SEEK1 protein
Consensus includes gb:R71245 /FEA=EST /DB_XREF=gi:844762 /DB_XREF=est:y54a05.s1 /CLONE=IMAGE:143072 /UG=Hs.174303 ESTs
mucin 3B
tubulin, beta, 5
hypothetical protein MGC10946
Consensus includes gb:AL080161.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434N144 (from clone DKFZp434N144) /FEA=mRNA /DB_XREF=gi:5262623 /UG=Hs.2104
chromodomain helicase DNA binding protein 4
small inducible cytokine subfamily A (Cys-Cys), member 19
Rac8p, a meiotic recombination and sister chromatid cohesion phosphoprotein of the rad21p family
hypothetical protein FLJ20274
hypothetical protein
Consensus includes gb:M14087.1 /DEF=Human HL14 gene encoding beta-galactoside-binding lectin, 3 end, clone 2 /FEA=mRNA /DB_XREF=gi:187113 /UG=Hs.287389
lymphotoxin beta receptor (TNFR superfamily, member 3)
pre-B-cell leukemia transcription factor 2
phosphodiesterase 4C, cAMP-specific (phosphodiesterase E1 dunce homolog, Drosophila)
plexin C1
urotensin 2
mexalonate kinase (mexalonate aciduria)
Consensus includes gb:S67777 /DEF=estrogen receptor (exon 6) human, tamoxifen-resistant breast tumor 17, Genomic Mutant, 187 nt /FEA=CDS_1 /DB_XREF=gi:45938: S67777
nurt-related transcription factor 1 (acute myeloid leukemia 1; ami1 oncogene)
development and differentiation enhancing factor 1
melanoma antigen, family B, 4
glucose phosphate isomerase
carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)
PRO0644 protein
solute carrier family 10 (sodium/bile acid cotransporter family), member 2
leukocyte membrane antigen
cytochrome P450, subfamily I (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile)
hypothetical protein FLJ14082
sex comb on midleg homolog 1
hypothetical protein DKFZp761C07121
CD28 antigen (Tp44)
melanoma antigen, family A, 10
Consensus includes gb:AK025276.1 /DEF=Homo sapiens cDNA: FLJ21623 fis, clone COL07915 /FEA=mRNA /DB_XREF=gi:0437759 /UG=Hs.306791 Homo sapiens cDNA
pre-B-cell leukemia transcription factor 2
hypothetical zinc finger protein FLJ14011
Consensus includes gb:AW298127 /FEA=EST /DB_XREF=gi:6704763 /DB_XREF=est:U1-H-BW0-ajs-e-05-Q-U1.s1 /CLONE=IMAGE:2733008 /UG=Hs.118632 ESTs, Highly
nuclear receptor coactivator 6 interacting protein

single-stranded DNA binding protein 2	NM_012446	1.33
hypothetical protein FLJ2282	NM_024792	1.33
paired basic amino acid cleaving system 4	D26514	1.33
melanoma antigen, family C, 1	AL022152	1.33
itchy homolog E3 ubiquitin protein ligase (mouse)	AB056663	1.33
Consensus includes gb:AV741657 /FEA=EST /DB_XREF=est:AV741657 /CLONE=CBMALG01 /UG=Hs.154085 leucine zipper protein 1	AV741657	1.33
Consensus includes gb:AK022170.1 /DEF=Homo sapiens cDNA FLJ12103 fls, clone MAMMA1000009 /FEA=mRNA /DB_XREF=gi:10433505 /UG=Hs.250813 Homo sapiens	AK022170	1.33
Putative prostate cancer tumor suppressor	AU158251	1.33
PP1201 protein	NM_022152	1.33
cholinergic receptor, nicotinic, alpha polypeptide 4	X89745	1.33
vesicle-associated membrane protein 4	BC005974	1.33
par-3 partitioning defective 3 homolog (C. elegans)	NM_019819	1.33
KIA0450 gene product	NM_014638	1.33
N-acetylglucosaminyl-1-phosphodiester alpha-N-acetylglucosaminidase	NM_016256	1.33
dihydropyrimidine S-succinyltransferase (E2 component of 2-oxo-glutarate complex)	AC006530	1.33
KIA00907 protein	NM_014949	1.33
mannosyl (alpha-1,3)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isoenzyme A	NM_012214	1.33
MHC class II transactivator	U18288	1.33
N-myc downstream regulated	NM_006096	1.33
sperm specific antigen 2	NM_006751	1.33
KIA00643 protein	AA207013	1.33
CGI-72 protein	NM_016018	1.33
recombinant binding protein suppressor of hairless (Drosophila)-like	NM_014276	1.33
protocadherin beta 8	NM_019120	1.33
DNA segment on chromosome 6(unique) 2654 expressed sequence	NM_012135	1.33
inhibin beta A (activin A, activin AB alpha polypeptide)	NM_002192	1.33
keratin, hair, acidic, 6	NM_003771	1.33
solute carrier family 18 (vesicular acetylcholine), member 3	NM_003055	1.33
complement-c1q tumor necrosis factor-related protein; likely ortholog of mouse CORS26 (collagenous repeat-containing sequence of 26-kDa protein)	NM_030945	1.33
death-associated protein kinase 2	NM_014328	1.33
basic helix-loop-helix domain containing, class B, 2	NM_003670	1.33
crystallin, mu	NM_001888	1.33
osteomodulin	NM_005014	1.33
CGI-105 protein	AA723370	1.33
28kD interferon responsive protein	NM_022147	1.33
hypothetical protein FLJ13215	NM_025004	1.33
folistatin	NM_006350	1.33
growth differentiation factor 5 (cartilage-derived morphogenetic protein-1)	NM_000557	1.33
bone morphogenetic protein receptor, type II (serine/threonine kinase)	NM_005110	1.33
fibulin 1	NM_006487	1.33
thymidine kinase 2, mitochondrial	BE895437	1.33
PRO0132 protein	NM_014116	1.33
solute carrier family 24 (sodium/potassium/calcium exchanger), member 1	AF026132	1.33
similar to rat integral membrane glycoprotein POM121	AU158490	1.33
keratin 6A	J00269	1.33
E2F-like protein	NM_016521	1.33
flavin containing monooxygenase 1	NM_002021	1.33
Consensus includes gb:AK024976.1 /DEF=Homo sapiens cDNA: FLJ21323 fls, clone COL02374, /FEA=mRNA /DB_XREF=gi:10437405 /UG=Hs.75914 Homo sapiens cDNA	AK024976	1.33
potassium channel, subfamily K, member 3 (TASK-1)	NM_002246	1.33
regulator of G-protein signalling 4	BC000737	1.33
signal transducer and activator of transcription 1, 91kD	NM_007315	1.33
G1 to S phase transition 2	NM_018094	1.33
solute carrier family 17 (sodium phosphate), member 3	NM_006632	1.33
fibromodulin	NM_002023	1.33
dom-3 homolog Z (C. elegans)	NM_005510	1.33
neurofilament, heavy polypeptide (200kD)	NM_021076	1.33
oxysterol-binding protein-related protein 1	W19983	1.33
Consensus includes gb:AK021614.1 /DEF=Homo sapiens cDNA FLJ11552 fls, clone HEMBA1003021, /FEA=mRNA /DB_XREF=gi:10432830 /UG=Hs.306600 Homo sapiens	AK021614	1.33
KIA0280 protein	D87470	1.33
calcium/calmodulin-dependent protein kinase kinase 2, beta	AK024748	1.33
a disintegrin and metalloproteinase domain 22	NM_004194	1.33
retinoic acid receptor, beta	NM_000965	1.33

prosudin
 KIAA0810 protein
 lipopolysaccharide binding protein
 solute carrier family 2 (facilitated glucose transporter), member 6
 epithelial membrane protein 1
 sodium channel, nonvoltage-gated 1, delta
 cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 18
 putative zinc finger protein from EUROMAGE 566589
 synaptophysin 3
 Consensus includes gb:AW516297 /FEA=EST /DB_XREF=gi:7154379 /DB_XREF=est:xd64d08.x1 /CLONE=IMAGE:2791215 /UG=Hs.323231 Homo sapiens cDNA FLJ111 AW516297
 hypothetical protein
 Consensus includes gb:BC545769 /FEA=EST /DB_XREF=gi:13544434 /DB_XREF=est:50257042F1 /CLONE=IMAGE:4701250 /UG=Hs.9927 Homo sapiens mRNA; cDN BG545769
 cyclin-dependent kinase inhibitor 1C (p57, Kip2)
 PTH-responsive osteosarcoma B1 protein
 purinergic receptor P2Y, G-protein coupled, 1
 facioscapular dysplasia (Aarskog-Scott syndrome)
 transducin-like enhancer of split 1 (E(spl1) homolog, Drosophila)
 hypothetical protein MGC10924 similar to Nedd4 WW-binding protein 5
 KIAA0551 protein
 protein phosphatase 1, regulatory (inhibitor) subunit 12B
 cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 8
 Consensus includes gb:AK023918.1 /DEF=Homo sapiens cDNA FLJ13656 fls, clone THYRO1000988 /FEA=mRNA /DB_XREF=gi:10436003 /UG=Hs.288489 Homo sapiens cDNA FLJ111 AW516297
 ras homolog gene family, member
 Consensus includes gb:AK024455.1 /DEF=Homo sapiens mRNA for FLJ00047 protein, partial cds. /FEA=mRNA /GEN=FLJ00047 /PROD=FLJ00047 protein /DB_XREF=gi:AK024455
 lipase, gastric
 hemopoietic cell kinase
 Consensus includes gb:AU148086 /FEA=EST /DB_XREF=gi:11009607 /DB_XREF=est:AU148086 /CLONE=MAMMA1002554 /UG=Hs.302681 Homo sapiens cDNA FLJ12 AU148086
 Consensus includes gb:AA707189 /FEA=EST /DB_XREF=gi:2717117 /DB_XREF=est:32g01.a1 /CLONE=IMAGE:452016 /UG=Hs.298998 Homo sapiens, Similar to hypot AA707189
 hypothetical protein FLJ11294
 crystallin leukodione CysLT2 receptor, cDNA; PSEC0146 from clone PLACE1006979
 chromosome X open reading frame 1
 A TP-binding cassette, sub-family A (ABC1), member 1
 eukaryotic translation initiation factor 2, alpha kinase 3
 PRO0628 protein
 T-cell lymphoma invasion and metastasis 2
 nuclear receptor co-repressor 2
 alpha2,8-sialyltransferase
 suppressor of K+ transport defect 1
 Consensus includes gb:AI186145 /FEA=EST /DB_XREF=gi:3736783 /DB_XREF=est:q646a06.x1 /CLONE=IMAGE:1742002 /UG=Hs.154338 Homo sapiens clone 24583 tr AI186145
 nuclear respiratory factor 1
 KIAA0878 protein
 sprouty homolog 1, antagonist of FGF signaling (Drosophila)
 protein phosphatase 1B (formerly 2C), magnesium-dependent, beta isoform
 aspartate beta-hydroxylase
 KIAA0638 protein
 F-box and WD-40 domain protein 1B
 regulator of G-protein signalling 6
 Consensus includes gb:AA527433 /FEA=EST /DB_XREF=gi:2269502 /DB_XREF=est:mg41b03.a1 /CLONE=IMAGE:337325 /UG=Hs.14846 Homo sapiens mRNA; cDNA D AA527433
 brefeldin A-inhibited guanine nucleotide-exchange protein 1
 interleukin 12B (natural killer cell stimulatory factor 2, cytotoxic lymphocyte maturation factor 2, p40)
 KIAA0775 gene product
 transforming growth factor, alpha
 stannocalcin 1
 insulin-like growth factor 1 receptor
 lymphocyte antigen 6 complex, locus E
 Consensus includes gb:BF47272 /FEA=EST /DB_XREF=gi:11512410 /DB_XREF=est:7m47a12.x1 /CLONE=IMAGE:3648695 /UG=Hs.288190 ESTs
 platelet-activating factor acetylhydrolase, isoform Ib, alpha subunit (45KD)
 lamin A/C
 angiotensinogen 2
 neurofilament, heavy polypeptide (200KD)
 Consensus includes gb:BG478428 /FEA=EST /DB_XREF=gi:13410807 /DB_XREF=est:602523839F1 /CLONE=IMAGE:4642353 /UG=Hs.326416 Homo sapiens mRNA; cDN BG478428
 laminin, beta 4
 AF028325
 carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)
 M76742

F-box and WD-40 domain protein 7 (archipelago homolog, Drosophila)
 proteasome (prosome, macropain) 26S subunit, ATPase, 4
 hypothetical protein LOC37147
 Consensus includes gb:BF000251 /FEA=EST /DB_XREF=gi:10700526 /DB_XREF=est:7h24e02.x1 /CLONE=IMAGE:3316922 /UG=Hs.227835 KIAA1049 protein
 putative transmembrane protein
 S-phase kinase-associated protein 2 (p45)
 Consensus includes gb:AW69655 /FEA=EST /DB_XREF=gi:8159499 /DB_XREF=est:EST381732 /UG=Hs.168386 ESTs
 photoblin-like protein MDS019
 interleukin 24
 lysyl oxidase-like 2
 Consensus includes gb:AU159040 /FEA=EST /DB_XREF=gi:11020561 /DB_XREF=est:AU159040 /CLONE=THYRO1000676 /UG=Hs.296745 Homo sapiens cDNA FLJ131
 Consensus includes gb:AW851559 /FEA=EST /DB_XREF=gi:7947076 /DB_XREF=est:IL3-CT0221-040400-107-E09 /UG=Hs.202695 Human soluble CD44 (CD44) mRNA, AW851559
 fibroblast growth factor 5
 hypothetical protein FLJ22344
 acid-inducible phosphoprotein
 hypothetical protein FLJ10147
 integrin, alpha M (complement component receptor 3, alpha; also known as CD11b (p170), macrophage antigen alpha polypeptide)
 KIAA1564 protein
 calbindin 1, (28KD)
 Consensus includes gb:AK024584.1 /DEF=Homo sapiens cDNA: FLJ20931 fls, done ADSE01282. /FEA=mRNA /DB_XREF=gi:10436896 /UG=Hs.203076 Homo sapiens c
 myeloid cell leukemia sequence 1 (BCL2-related)
 Raf2 interacting protein 8
 NG3 protein
 Consensus includes gb:A921841 /FEA=EST /DB_XREF=gi:5657805 /DB_XREF=est:wp07a09.x1 /CLONE=IMAGE:2464120 /UG=Hs.212670 ESTs
 guanylate cyclase 2f, retinal
 GATA binding protein 6
 hypothetical protein MGCS287
 epidermal growth factor receptor substrate EPS15R
 Consensus includes gb:BF348061 /FEA=EST /DB_XREF=gi:112956556 /DB_XREF=est:602022810F1 /CLONE=IMAGE:4157907 /UG=Hs.167988 neural cell adhesion mole
 KIAA0461 protein
 catenin (cadherin-associated protein), alpha 1 (102KD)
 hypothetical protein FLJ11580
 cullin 3
 BH3 interacting domain death agonist
 papillomavirus regulatory factor PRF-1
 B melanoma antigen
 KIAA0444 protein
 epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)
 cytosolic ovarian carcinoma antigen 1
 1-acylglycerol-3-phosphate O-acyltransferase 3
 RNA binding motif protein 12
 retinoblastoma binding protein 4
 KIAA0015 gene product
 Consensus includes gb:U63332.1 /DEF=Human super cysteine rich protein mRNA, partial cds. /FEA=mRNA /PROD=super cysteine rich protein /DB_XREF=gi:1480862 /U
 sodium channel, voltage-gated, type II, beta polypeptide
 KIAA1456 protein
 SWI/SNF related, actin dependent regulator of chromatin, subfamily d, member 3
 Consensus includes gb:M31949 /DEF=Human Ig rearranged mu-chain V-region gene, subgroup VH-III, exon 1 and 2 /FEA=CDS /DB_XREF=gi:185254 /UG=Hs.247923 Hs
 myelin transcription factor 1-like
 retinoblastoma binding protein 9
 mannosidase, alpha, class 2A, member 2
 Consensus includes gb:AK024123.1 /DEF=Homo sapiens cDNA FLJ14061 fls, clone HEMBB 1000749, /FEA=mRNA /DB_XREF=gi:10436427 /UG=Hs.306684 Homo sapie
 dedicator of cyto-kinesis 1
 Consensus includes gb:BG025063 /FEA=EST /DB_XREF=gi:12411278 /DB_XREF=est:602276408F1 /CLONE=IMAGE:4364304 /UG=Hs.44888 ESTs
 NK3 transcription factor homolog A (Drosophila)
 prospero-related homeobox 1
 neuraxin 2
 E1B-55kDa-associated protein 5
 KIAA0300 protein
 chromosome 21 open reading frame 80
 protease, serine, 3 (trypsin 3)
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily b, member 1
 Consensus includes gb:AC004862 /DEF=Homo sapiens PAC clone RP4-697H17 from Tq11.23-q21.1 /FEA=CDS /DB_XREF=gi:3900856 /UG=Hs.247768 Homo sapiens P AC004862

proteasome (prosome, macropain) subunit, beta type, 1	W88293	1.32
prolactin receptor	S78505	1.32
unit-related transcription factor 2	NM_004348	1.32
RUB-4b protein	NM_016530	1.32
leucine zipper, putative tumor suppressor 1	AF123854	1.32
Consensus includes gb:AL035406 /DEF=Human DNA sequence from clone RP1-233K16 on chromosome 1p36.21-36.33 Contains the gene KIAA0444, a putative chromod	AL035406	1.32
likely ortholog of mouse synaptic vesicle glycoprotein 2a	NM_014849	1.32
phosphatidyserine synthase 2	NM_030783	1.32
hypothetical protein FLJ22601	NM_024822	1.32
arachidonate 5-lipoxygenase	NM_000698	1.32
NPHS2 gene (podocin)	NM_014825	1.32
dead ringer (Drosophila)-like 2 (bright and dead ringer)	NM_008465	1.32
type 1 protein phosphatase inhibitor	NM_025210	1.32
G protein-coupled receptor 49	AF062006	1.32
paraneoplastic cancer-testis-brain antigen	NM_013364	1.32
taste receptor, type 2, member 10	NM_023921	1.32
ubiquitin-conjugating enzyme E2G 1 (UBC7 homolog, C. elegans)	AW299555	1.32
hypothetical protein FLJ12765	NM_024852	1.32
killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2	NM_006737	1.32
hypothetical protein FLJ10381	NM_018080	1.32
mannose-6-phosphate receptor (cation dependent)	A583537	1.32
hypothetical protein FLJ14351	NM_024732	1.32
TBC1 (tre-2/USP6, BUR2, cdc16) domain family, member 1	BE875153	1.32
hypothetical protein FLJ20581	A733019	1.32
sarcoma antigen	NM_016666	1.32
Consensus includes gb:U81961 /DEF=Homo sapiens amiloride-sensitive epithelial sodium channel alpha subunit alpha ENaC-1 and alpha ENaC-2 (SCNN1A) genes, allert	U81961	1.32
solute carrier family 8 (sodium/calcium exchanger), member 1	AF106389	1.32
brain cell membrane protein 1	A803181	1.32
CGI-83 protein	NM_016027	1.32
Consensus includes gb:AA446466 /FEA=EST /DB_XREF=gi:2159131 /DB_XREF=est:zw65b12.s1 /CLONE=IMAGE:781055 /UG=Hs.193783 Human DNA sequence from c	AA446466	1.32
actin filament associated protein	NM_021638	1.32
v-ets erythroblastosis virus E26 oncogene homolog 2 (avian)	AL575509	1.32
5-methyltetrahydrofolate-homocysteine methyltransferase reductase	NM_024010	1.32
actinin, alpha 2	W63731	1.32
aldehyde dehydrogenase 3 family, member A2	L47162	1.32
G protein-coupled receptor 21	AF129166	1.32
Consensus includes gb:AF088790.1 /DEF=Homo sapiens aconitase precursor (ACON) mRNA, nuclear gene encoding mitochondrial protein, partial cds. /FEA=mRNA /GEI	NM_005294	1.32
Consensus includes gb:AL049767 /DEF=Human DNA sequence from clone 172H20 on chromosome 20q12-13.12 Contains genes for serpinogelin I and II and PI3 (proteas	AF088790	1.32
Consensus includes gb:Z83929 /DEF=Human DNA sequence from clone 272E8 on chromosome Xp22.13-22.31. Contains a pseudogene similar to MDM2-Like P53-binding	AL049767	1.32
keratin, hair, basic, 3	Z83929	1.32
hepatocyte nuclear factor 4, alpha	NM_002282	1.32
nuclear protein, marker for differentiated aortic smooth muscle and down-regulated with vascular injury	X87870	1.32
phospholipase A2, group VI (cytosolic, calcium-independent)	NM_005876	1.32
neuropilin 1	AF102988	1.32
hypothetical protein FLJ14031	AF280547	1.32
mitochondrial ribosomal protein L9	AK024083	1.32
chloride channel 7	AF225421	1.32
similar to rat myomegalin	AW611641	1.32
flavin containing monooxygenase 4	Z67743	1.32
integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3 receptor)	NM_022169	1.32
NADPH-dependent FMN and FAD containing oxidoreductase	BC005849	1.32
cadherin 11, type 2, OB-cadherin (osteoblast)	NM_002022	1.32
nuclear receptor subfamily 4, group A, member 3	NM_002204	1.32
vitamin D (1,25-dihydroxyvitamin D3) receptor	NM_014434	1.32
AP5-like 1	NM_001787	1.32
hypothetical protein FLJ11259	NM_006981	1.32
SH3-domain GRB2-like 3	AAA45701	1.32
hypothetical protein FLJ10244	NM_018370	1.32
interleukin 3 receptor, alpha (low affinity)	AF036272	1.32
postmeiotic segregation increased 2-like 4	NM_018037	1.32
	NM_002183	1.32
	D38500	1.32

a disintegrin and metalloproteinase domain 12 (meltrin alpha)
U1-snRNP binding protein homolog (70kD)
translocase of inner mitochondrial membrane 17 homolog A (yeast)
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13
syncytin, alpha (non A4 component of amyloid precursor)
tomosyn
Interferon-induced protein 35
immunoglobulin heavy constant mu
PRO1768 protein
potassium inwardly-rectifying channel, subfamily J, member 2
mitogen-activated protein kinase 8 interacting protein 1
Consensus includes gb:AL109809 /DEF=Human DNA sequence from clone RP4-873D20 on chromosome 20. Contains a PTPNS (protein tyrosine phosphatase, non-recap AL109809
Consensus includes gb:BE748563 /FEA=EST /DB_XREF=gi:10162555 /DB_XREF=est:60157152111 /CLONE=IMAGE:3838541 /UG=Hs.13413 Homo sapiens clone 2446: BE748563
makorin, ring finger protein, 2
cytochrome P450, subfamily IIE (ethanol-inducible)
collagen, type IV, alpha 3 (Goodpasture antigen)
hypothetical protein FLJ11383
prolaserone (prosome, macropain) 26S subunit, ATPase, 2
Consensus includes gb:AF070566.1 /DEF=Homo sapiens clones 24611 and 24705 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387934 /UG=Hs.302042 Homo sapiens
N-acetylglucosaminidase, alpha- (Sanfilippo disease IIIB)
postmeiotic segregation increased 2-like 3
Dombrock blood group
dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)
integrin, beta 5
Consensus includes gb:AF003737.1 /DEF=Homo sapiens chromosome 21q22.1 anonymous mRNA sequence. /FEA=mRNA /DB_XREF=gi:2197088 /UG=Hs.302179 Hom AF003737
gb:AF130057.1 /DEF=Homo sapiens clone FL85539 PRO1454 mRNA, complete cds. /FEA=mRNA /PROD=PRO1454 /DB_XREF=gi:11493420 /UG=Hs.323288 Homo sap AF130057
hypothetical protein FLJ20489
Jun D proto-oncogene
hypothetical protein FLJ20584
Consensus includes gb:BE044440 /FEA=EST /DB_XREF=gi:8361493 /DB_XREF=est:th045a07.x1 /CLONE=IMAGE:3040356 /UG=Hs.251278 KIAA1201 protein
KIAA1024 protein
immunoglobulin heavy constant gamma 3 (G3m marker)
apolipoprotein L, 5
pervin, beta
serma domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F
Consensus includes gb:H95263 /FEA=EST /DB_XREF=gi:1102896 /DB_XREF=est:W20604.s1 /CLONE=IMAGE:234366 /UG=Hs.13431 Homo sapiens clone 24790 mRN H95263
bromodomain-containing 2
small protein effector 1 of Cdc42
Interleukin 17E
transmembrane, prostate androgen induced RNA
adenylate kinase 1
Consensus includes gb:AW405975 /FEA=EST /DB_XREF=gi:6925032 /DB_XREF=est:U1-HF-BL0-acc-a-05-0-U1.r1 /CLONE=IMAGE:3060366 /UG=Hs.247721 Homo saple AW405975
podocalyxin-like
glutathione peroxidase 3 (plasma)
hypothetical protein FLJ20668
adaptor-related protein complex 1, mu 2 subunit
adenosine A1 receptor
bladder cancer overexpressed protein
ATPase, H⁺ transporting, lysosomal (vacuolar proton pump) 9kD
hemochromatosis
hypothetical protein FLJ11220
taste receptor, type 2, member 13
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 31
endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 4
melanoma antigen, family A, 6
mas-21-like 2 (C. elegans)
replication protein A complex 34 kd subunit homolog Rpa4
inositol 1,4,5-bisphosphate receptor, type 1
hypothetical protein SBB187
protein phosphatase 1, regulatory (inhibitor) subunit 12B
hypothetical protein FLJ20701
Bacillus subtilis /REF=X04603 /LEN=2073 (-5, -M, -3 represent transcript regions 5 prime X04603
guanine nucleotide binding protein 11

gb:AF343666.1 /DEF=Homo sapiens translocation associated fusion protein IRTA1IGA1 (IRTA1IGA1) mRNA, complete cds. /FEA=mRNA /GEN=IRTA1IGA1 /PROD=tr AF343666
 uridine monophosphate synthetase (uridine phosphorylase) transferase and oridine-5'-decarboxylase)
 actin like protein
 olfactory receptor, family 7, subfamily C, member 1
 dipeptidylpeptidase VI
 keratin 4
 Consensus includes gb:AK023345.1 /DEF=Homo sapiens cDNA FLJ13283 fs, clone OVARC1001113, highly similar to Homo sapiens diaphanous 1 (HDIA1) mRNA. /FEA=AK023345
 Consensus includes gb:AK023345.1 /DEF=Homo sapiens cDNA FLJ13283 fs, clone OVARC1001113, highly similar to Homo sapiens diaphanous 1 (HDIA1) mRNA. /FEA=AK023345
 alkaline phosphatase, placental (Regan isozyme)
 hypothetical protein FLJ21522
 phosphoglucomutase 5
 Consensus includes gb:AK025026.1 /DEF=Homo sapiens cDNA: FLJ21373 fs, clone COL03214. /FEA=mRNA /DB_XREF=gi:10437461 /UG=Hs.306747 Homo sapiens cD AK025026
 protein phosphatase 1, regulatory (inhibitor) subunit 14D
 hypothetical protein FLJ10698
 prostaglandin-endoperoxide synthase 2 (prostaglandin GH synthase and cyclooxygenase)
 BTG3 associated nuclear protein
 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3
 hypothetical protein FLJ22814
 4-hydroxyphenylpyruvate dioxygenase
 synapsin II
 NK3 transcription factor homolog A (Drosophila)
 PCTAIRE protein kinase 3
 POU domain, class 4, transcription factor 2
 DNA fragmentation factor, 45 kD, alpha polypeptide
 a disintegrin and metalloproteinase domain 20
 polycythemia rubra vera 1; cell surface receptor
 hypothetical protein PRO1817
 Huntingtin interacting protein B
 sphingomyelin phosphodiesterase 2, neutral membrane (neutral sphingomyelinase)
 latrophilin
 killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 2
 CUG triplet repeat, RNA binding protein 2
 nitric oxide synthase 3 (endothelial cell)
 DKFZP564B0769 protein
 egi nine homolog 3 (C. elegans)
 solute carrier family 14 (urea transporter), member 2
 transferrin
 hypothetical protein FLJ10375
 Consensus includes gb:AK026155.1 /DEF=Homo sapiens cDNA: FLJ22502 fs, clone HRC11383. /FEA=mRNA /DB_XREF=gi:10438915 /UG=Hs.298140 Homo sapiens cD AK026155
 acyl-Coenzyme A dehydrogenase, long chain
 plasminogen activator, urokinase receptor
 ATP-binding cassette, sub-family C (CFTR/MRP), member 6
 Consensus includes gb:AL162044.1 /DEF=Homo sapiens mRNA: cDNA DKFZp761L0812 (from clone DKFZp761L0812); partial cds. /FEA=mRNA /GEN=DKFZp761L0812 AL162044
 Consensus includes gb:AL162044.1 /DEF=Homo sapiens mRNA: cDNA DKFZp761L0812 (from clone DKFZp761L0812); partial cds. /FEA=mRNA /GEN=DKFZp761L0812 AL162044
 KIAA0551 protein
 Consensus includes gb:AF007137.1 /DEF=Homo sapiens clone 23618 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852612 /UG=Hs.302185 Homo sapiens clone 23618 AF007137
 pregnancy-associated plasma protein A
 DKFZP586P0123 protein
 p632 related 1
 capicua homolog (Drosophila)
 HT021
 Huntingtin interacting protein M
 G-protein coupled receptor
 potassium voltage-gated channel, shaker-related subfamily, member 2
 Werner helicase interacting protein
 crystallin, gamma A
 leptin receptor
 geranylgeranyl diphosphate synthase 1
 Rls
 HEAT-like repeat-containing protein
 crystallin, beta B1
 engrailed homolog 1
 ADP-ribosylation factor-like 5
 Consensus includes gb:BF222018 /FEA=EST /DB_XREF=gi:111129195 /DB_XREF=ast:7p41a04.x1 /CLONE=IMAGE:364827 /UG=Hs.289803 ESTs. Moderately similar to BF222018

a disintegrin and metalloproteinase domain 22	AF155381	1.31
carbohydrate (chondroitin 6/keratan) sulfotransferase 3	AB017915	1.31
sepin 6	AK026589	1.31
microfibrillar-associated protein 3	R38475	1.31
solute carrier family 25 (mitochondrial carrier; ornithine transporter) member 15	NM_014252	1.31
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4	BF223021	1.31
KIAA0716 gene product	NM_014705	1.31
phosphatidic acid phosphatase type 2A	AB000888	1.31
integrin, beta 5	NM_002213	1.31
Consensus includes gb-AK439416 /FEA=EST /DB_XREF=gi:4304169 /DB_XREF=est:ij04b04.x1 /CLONE=IMAGE:2140495 /UG=Hs.165728 ESTs	AI439416	1.31
Rho GTPase activating protein 6	NM_006125	1.31
alkaline phosphatase, liver/bone/kidney	X14174	1.31
discolidin domain receptor family, member 1	L11315	1.31
NPC1 (Niemann-Pick disease, type C1, gene)-like 1	NM_013389	1.31
integrin, beta 3 (platelet glycoprotein IIb, antigen CD61)	U95204	1.31
solute carrier family 4, sodium bicarbonate cotransporter, member 4	NM_003759	1.31
E3 ubiquitin ligase SMURF1	AF199384	1.31
catalase	NM_001752	1.31
collapsin response mediator protein 1	NM_001313	1.31
arginase, type II	NM_001172	1.31
SMAD in the antisense orientation	NM_022001	1.31
Consensus includes gb-AW851559 /FEA=EST /DB_XREF=gi:7947078 /DB_XREF=est:il3-C70221-040400-107-E09 /UG=Hs.202695 Human soluble CD44 (CD44) mRNA, AW851559	AW851559	1.31
Consensus includes gb-A1810380 /FEA=EST /DB_XREF=gi:5386948 /DB_XREF=est:wb87h10.x1 /CLONE=IMAGE:2312707 /UG=Hs.47274 Homo sapiens mRNA, cDNA C A1810380	C A1810380	1.31
Consensus includes gb-BE464789 /FEA=EST /DB_XREF=gi:9510493 /DB_XREF=est:ch837a06.x1 /CLONE=IMAGE:3144178 /UG=Hs.27342 ESTs	BE464789	1.31
Consensus includes gb-AL109728.1 /DEF=Homo sapiens mRNA (full length insert cDNA clone EUROMAGE 124430) /FEA=EST /DB_XREF=gi:5689792 /UG=Hs.187513 AL109726	AL109726	1.31
hypothetical protein similar to CG7843	AL138752	1.31
solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	D50402	1.31
interleukin 1 receptor, type II	NM_004633	1.31
calpastatin	D16217	1.31
FKBP-associated protein	NM_007070	1.31
midline 1 (Opitz/BBB syndrome)	AF041209	1.31
Consensus includes gb-H49077 /FEA=EST /DB_XREF=gi:988918 /DB_XREF=est:yq18a05.s1 /CLONE=IMAGE:274180 /UG=Hs.159179 Homo sapiens mRNA sequence, I H49077	I H49077	1.31
hypothetical protein FLJ12681	A003763	1.31
adenylate kinase 3	NM_013410	1.31
diacyl, large (Drosophila) homolog 3 (neuroendocrine-dlg)	NM_021120	1.31
poreye protein 3	NM_022361	1.31
gb-BC002792.1 /DEF=Homo sapiens, clone MGC:3963, mRNA, complete cds. /FEA=EST /DB_XREF=gi:12803890 /UG=Hs.81 BC002792	BC002792	1.31
S184 protein	AV757384	1.31
neutrophilic tyrosine kinase, receptor, type 3	S76475	1.31
homeo box D1	NM_024501	1.31
KIAA1886 protein	AL136572	1.31
Consensus includes gb-AK026847.1 /DEF=Homo sapiens cDNA: FLJ23194 fs, clone REC00490. /FEA=EST /DB_XREF=gi:10439802 /UG=Hs.306887 Homo sapiens cDNA, AK026847	AK026847	1.31
chorionic somatomammotropin hormone 1 (placental lactogen)	NM_022640	1.31
activated leucocyte cell adhesion molecule	AA158721	1.31
ectodermal dysplasia 1, anhidrotic	AF061193	1.31
leucyl-RNA synthetase, mitochondrial	NM_015340	1.31
aldehyde dehydrogenase 3 family, member A1	NM_000691	1.31
acetylserotonin O-methyltransferase	BC001620	1.31
transcriptional activator of the c-fos promoter	NM_006365	1.31
Sp4 transcription factor	NM_003112	1.31
chromosome 4 open reading frame 5	BF939176	1.31
eyes absent homolog 2 (Drosophila)	U71207	1.31
tissue inhibitor of metalloproteinase 4	NM_003256	1.31
zinc finger protein 331; zinc finger protein 463	NM_018555	1.31
testis-specific transcript, Y-linked 2	AF000991	1.31
stem cell growth factor, lymphocyte secreted C-type lectin	D86586	1.31
PRO0246 protein	NM_014123	1.31
CGI-119 protein	NM_016056	1.31
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide, Y chromosome	AF000985	1.31
interferon, alpha 18	NM_002173	1.31
deoxyribonuclease I-like 1	NM_006730	1.31
Consensus includes gb-BE738425 /FEA=EST /DB_XREF=gi:10152417 /DB_XREF=est:50157244.T1 /CLONE=IMAGE:3839147 /UG=Hs.241507 ribosomal protein S6	BE738425	1.31
biliverdin reductase A	U34877	1.31

sema domain, immunoglobulin domain (Ig), and GPI membrane anchor, (semaphorin) 7A
 SAM domain, SH3 domain end nuclear localisation signals, 1
 GATA binding protein 2
 thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)
 proteasome (prosome, macropain) subunit, beta type, 2
 KIAA1199 protein
 basic-helix-loop-helix-PAS protein
 caspase 1, apoptosis-related cysteine protease (interleukin 1, beta, convertase)
 leucine-rich, glioma inactivated, 1
 glial cells missing homolog a (Drosophila)
 matrix metalloproteinase 20 (matrilysin)
 leucine-rich repeat-containing 3
 tumor necrosis factor receptor superfamily, member 10b
 catenin (cadherin-associated protein), alpha 1 (102kD)
 XAGE-1 protein
 hypothetical protein FLJ10922
 cellular retinoic acid binding protein 1
 duodenal cytochrome b
 macrophage scavenger receptor 1
 calyculin-2
 dynabrevin, beta
 Consensus includes gb:A1808793 /FEA=EST /DB_XREF=gi:5393359 /DB_XREF=est:wt15d05.x1 /CLONE=IMAGE:2350665 /UG=Hs.249239 collagen, type VIII, alpha 2
 fibroblast growth factor 14
 KIAA0530 protein
 hypothetical protein PRO1097
 aquaporin 4
 adipose specific 2
 DKFZP564O243 protein
 N-acetylated alpha-linked acidic dipeptidase 2
 villin 2 (ezrin)
 goliath protein
 calcium binding protein P22
 hypothetical protein FLJ20260
 RAB interacting factor
 Consensus includes gb:S65761.1 /DEF=anti-colorectal carcinoma heavy chain=glycoprotein CANAG-50 specific IgG1 kappa human, 19.9 hybridoma, antibody 1116NS19.9 S65761
 CDK2-associated protein 1
 forkhead box 11
 PRO1892 protein
 multiple PDZ domain protein
 reversion-inducing-cysteine-rich protein with kazal motifs
 myogenin (myogenic factor 4)
 Consensus includes gb:AF090910.1 /DEF=Homo sapiens clone HQ0262. /FEA=mRNA /DB_XREF=gi:6690195 /UG=Hs.236760 Homo sapiens clone HQ0262
 glutamate receptor, ionotropic, kainate 1
 vacuolar protein sorting 11 (yeast)
 vesicle-associated membrane protein 2 (synaptobrevin 2)
 UDP-Gal4betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 3
 Consensus includes gb:AL121890 /DEF=Human DNA sequence from clone RPS-1116H23 on chromosome 20 Contains a novel gene, a 40S ribosomal protein S21 pseudo
 keratin associated protein 8.9
 KIAA1841 protein
 hepatitis A virus cellular receptor 1
 DKFZP568A0522 protein
 Consensus includes gb:AW954107 /FEA=EST /DB_XREF=gi:8143790 /DB_XREF=est:EST366177 /UG=Hs.303394 Homo sapiens cDNA: FLJ22150 f1s, clone HRC00108
 MKP-1 like protein tyrosine phosphatase
 lymphoid blast crisis oncogene
 phosphoinositide-3-kinase, catalytic, beta polypeptide
 lysozyme, alpha-L
 sprouty (Drosophila) homolog 4
 casein kinase 1, gamma 2
 Consensus includes gb:AK073448.1 /DEF=Homo sapiens cDNA FLJ13384 f1s, clone PLACE1001082, highly similar to Homo sapiens mRNA for lysine-ketoglutarate reductase
 Consensus includes gb:AL122074.1 /DEF=Homo sapiens mRNA, cDNA DKFZP434N0435 (from clone DN7ZP434N0435), partial cds. /FEA=mRNA /GEN=DKFZP434N0435
 KIAA1856 protein
 ubiquitin specific protease 9, Y chromosome (Drosophila fat facets related)
 KIAA0453 protein

AF071542 1.31
 NM_022136 1.31
 AL563460 1.31
 M24899 1.31
 BC000268 1.31
 AB033025 1.31
 NM_022123 1.31
 M87507 1.31
 NM_005097 1.31
 AB026483 1.31
 NM_004771 1.31
 NM_030891 1.31
 BC001281 1.31
 D14705 1.31
 NM_020411 1.31
 NM_018273 1.31
 NM_004378 1.31
 NM_024843 1.31
 A1299239 1.31
 NM_022131 1.31
 A1872185 1.31
 A1806793 1.31
 NM_004115 1.31
 BF223237 1.31
 NM_018499 1.31
 D63412 1.31
 NM_006829 1.31
 BC002571 1.31
 AJ012370 1.31
 AF189015 1.31
 AW194818 1.31
 NM_007236 1.31
 NM_017731 1.31
 U74324 1.31
 NM_004642 1.31
 NM_012188 1.31
 AF118086 1.31
 NM_003829 1.31
 AK022887 1.31
 NM_002479 1.31
 AF090910 1.31
 U16125 1.31
 NM_021729 1.31
 NM_014232 1.31
 AB050855 1.31
 AL121890 1.31
 NM_030875 1.31
 AB046881 1.31
 NM_012206 1.31
 AF113007 1.31
 AW954107 1.31
 NM_007026 1.31
 AF126008 1.31
 BC005393 1.31
 A142444.1 1.31
 NM_030964 1.31
 AL530441 1.31
 AK023446 1.31
 AL122074 1.31
 BE552334 1.30
 NM_004654 1.30
 AA524553 1.30

hypothetical protein FLJ10300
 biliverdin reductase A
 Consensus includes gb:AK025026.1 /DEF=Homo sapiens cDNA: FLJ21373 fls. clone COL03214. /FEA=mrRNA /DB_XREF=gi:10437461 /UG=Hs.306747 Homo sapiens cC
 melatonin receptor 1A
 KIAA1330 protein
 p21(CDKN1A)-activated kinase 6
 proteasome (prosome, macropain) 26S subunit, non-ATPase, 10
 glucagon-like peptide 1 receptor
 carbonyl reductase 3
 nuclear autoantigen
 hypothetical protein PRO1905
 cyclin-dependent kinase 5, regulatory subunit 1 (p35)
 homogenisate 1,2-dioxygenase (homogenisate oxidase)
 KIAA1218 protein
 galanin receptor 1
 opioid receptor, mu 1
 GCIP-interacting protein p29
 Consensus includes gb:AY295387 /FEA=EST /DB_XREF=gi:5702003 /DB_XREF=est:U1-H-B12-shv-e-04-O-U.s1 /CLONE=IMAGE:2728039 /UG=Hs.196585 ESTs, Weakt
 uromodulin (urmuocoid, Tamm-Horsfall glycoprotein)
 Consensus includes gb:AL049244.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564C163. /FEA=mrRNA /DB_XREF=gi:4498979 /UG=Hs.3070 AL049244
 Consensus includes gb:AK025650.1 /DEF=Homo sapiens cDNA: FLJ21997 fls. clone HEP06590. /FEA=mrRNA /DB_XREF=gi:10438234 /UG=Hs.306820 Homo sapiens cD
 mitochondrial NAD+-dependent retinol dehydrogenase 4
 biliverdin reductase A
 H factor 1 (complement)
 PHD finger protein 2
 cyclin-dependent kinase 3
 KIAA0721 protein
 KIAA0605 gene product
 adaptor-related protein complex 1, gamma 1 subunit
 v-aki sarcoma viral oncogene homolog (avian)
 arctin silent mating type information regulation 2 homolog 1 (S. cerevisiae)
 alpha-1-microglobulin/bikunin precursor
 axonal transport of synaptic vesicles
 runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)
 likely ortholog of mouse testis expressed gene 27
 epoxide hydrolase 1, microsomal (xenobiotic)
 hepatocellular carcinoma-associated antigen 127
 zinc finger protein 81 (HFZ20)
 insulin-like growth factor binding protein 3
 KIAA1193 protein
 hypothetical protein FLJ11561
 corticotropin releasing hormone receptor 2
 ADP-ribosylation factor-like 1
 lymphocyte-activation gene 3
 small inducible cytokine subfamily A (Cys-Cys), member 27
 hypothetical protein FLJ21079
 microtubule-associated protein 1A
 hypothetical protein FLJ10188
 ataxia telangiectasia and Rad3 related
 solute carrier family 8 (sodium/calcium exchanger), member 1
 ecdonucleotide pyrophosphatase/phosphodiesterase 3
 homeo box A1
 methionyl aminopeptidase 1
 prostaglandin E receptor 3 (subtype EP3)
 myosin IB
 Consensus includes gb:AU157078 /FEA=EST /DB_XREF=gi:11016599 /DB_XREF=est:AU157078 /CLONE=PLACE1006017 /UG=Hs.287580 Homo sapiens cDNA FLJ135
 metastasis-associated 1-like 1
 BRF1 homolog, subunit of RNA polymerase III transcription initiation factor IIIB (S. cerevisiae)
 hypothetical protein FLJ13480
 alpha2,3-sialyltransferase
 nuclear receptor subfamily 2, group F, member 6
 hexosaminidase B (beta polypeptide)

NM_018051
 NM_000712
 AK025028
 NM_005958
 AB037751
 AJ236915
 NM_002814
 L23503
 NM_001236
 AF243424
 NM_018608
 AL567411
 NM_000187
 AI017382
 NM_001480
 L29301
 NM_015484
 AW295387
 NM_003361
 AL050331
 NM_014694
 AL050025
 BF431501
 NM_012238
 AF019214
 AV645756
 BC647330
 D89788
 NM_021943
 NM_000120
 NM_018684
 X68011
 M31159
 AB033019
 NM_024651
 AF019381
 AU161560
 NM_002286
 NM_006664
 NM_024576
 AF095910
 NM_018017
 U49844
 AF128524
 NM_005021
 S76910
 D42084
 NM_000957
 BF432550
 NM_004739
 NM_001519
 NM_025120
 AB069587
 BC002669
 NM_000521

G protein-coupled receptor 42
 GLO2 protein
 pre-B-cell leukemia transcription factor 2
 Consensus includes gb:BF056029 /FEA=EST /DB_XREF=gi:10809925 /DB_XREF=est:h0703.x1 /CLONE=IMAGE:3443838 /UG=Ha.174910 ESTs. Weakly similar to cyst BF056029
 hypophthalic protein MG3156
 phospholipid scramblase 1
 frizzled homolog 7 (Drosophila)
 zinc finger protein RINZF
 gap junction protein, beta 3, 31kD (connexin 31)
 myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 4
 lysyl oxidase-like 2
 bone morphogenetic protein 5
 tropomodulin
 death-associated protein kinase 1
 protein kinase H11
 proteasome (prosome, macropain) subunit, alpha type, 7
 SH3-domain binding protein 4
 dynein, cytoplasmic, light intermediate polypeptide 2
 neural cell adhesion molecule 1
 centrosomal protein 2
 Down syndrome critical region gene 4
 neurogenin 3
 Homer, neuronal immediate early gene, 3
 zinc finger protein 202
 taste receptor, type 2, member 1
 a disintegrin and metalloproteinase domain 12 (meltrin alpha)
 hypophthalic protein FLJ11264
 KIAA0820 protein
 UL16 binding protein 2
 hypophthalic protein FLJ11952
 protein tyrosine phosphatase, receptor type, H
 hypophthalic protein FLJ11585
 trnucleotide repeat containing 11 (THR-associated protein, 230 kDa subunit)
 double C2-like domains, alpha
 interleukin 1, epsilon
 extracellular link domain-containing 1
 Consensus includes gb:AA631242 /FEA=EST /DB_XREF=gi:2553853 /DB_XREF=est:nq81b04.s1 /CLONE=IMAGE:1158703 /UG=Hs.21349 ESTs. Moderately similar to R
 uncharacterized gastric protein ZAS2P
 arachne homolog 2 (Drosophila)
 Interleukin 1 receptor antagonist
 Rag D protein
 solute carrier family 2 (facilitated glucose transporter), member 3
 homeo box D9
 zinc finger protein 180 (HZF168)
 killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 3
 opsin 3 (enkephalopsin, panopsin)
 Consensus includes gb:BE465829 /FEA=EST /DB_XREF=gi:9511604 /DB_XREF=est:h080n09.x1 /CLONE=IMAGE:3194177 /UG=Hs.150539 Homo sapiens cDNA FLJ137
 A kinase (PRKA) anchor protein 7
 uncoupling protein 3 (mitochondrial, proton carrier)
 synapjanin 2
 Consensus includes gb:AF229166.1 /DEF=Homo sapiens docking protein 1-like protein mRNA, partial cds. /FEA=mRNA /PROD=docking protein 1-like protein /DB_XREF=AF229166
 NT-REN-45 antigen
 corticotropin releasing hormone receptor 1
 hypophthalic protein DKFZp434N1923
 UDP-N-acetyl-alpha-D-galactosamine polypeptide N-acetylglucosaminyltransferase 3 (GlcNAc-T3)
 olfactory receptor, family 1, subfamily D, member 4
 contactin associated protein-like 2
 kininogen
 Consensus includes gb:AL137624.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M1812 (from clone DKFZp434M1812). /FEA=mRNA /DB_XREF=gi:8808420 /UG=Hs.3C /AL137624
 B9 protein
 small nuclear ribonucleoprotein polypeptide F
 SH3-domain binding protein 1
 nucleosome assembly protein 1-like 1

NM_005305
 1.30
 NM_020183
 1.30
 NM_002586
 1.30
 Consensus includes gb:BF056029 /FEA=EST /DB_XREF=gi:10809925 /DB_XREF=est:h0703.x1 /CLONE=IMAGE:3443838 /UG=Ha.174910 ESTs. Weakly similar to cyst BF056029
 1.30
 NM_024077
 1.30
 NM_021105
 1.30
 AI333651
 1.30
 NM_023929
 1.30
 AF098730
 1.30
 AB011358
 1.30
 AF117949
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 NM_021073
 1.30
 NM_003275
 1.30
 BC003614
 1.30
 AF133207
 1.30
 NM_002792
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 AF015043
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 AB040861
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 M22094
 1.30
 BC001433
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 NM_005867
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 NM_020999
 1.30
 AB71287
 1.30
 N46430
 1.30
 NM_016599
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 NM_021841
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 NM_018371
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 AL136712
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 NM_025217
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 AK022014
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 NM_002842
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 BF476502
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 AI217362
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 NM_003586
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 NM_014440
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 AL574194
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 AA631242
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 NM_020669
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 AV694434
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 AW083357
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 AL136717
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 AA778684
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 NM_014213
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 NM_013256
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 NM_012313
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 NM_014322
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 BE465829
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 AF161075
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 NM_003356
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 AK026758
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 AF229166
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 NM_016121
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 X72304
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 BF063271
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 NM_003552
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 NM_014141
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 NM_000893
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 NM_00137624
 1.30
 NM_015681
 1.30
 NM_003095
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 NM_018957
 1.30
 BC002387
 1.30

pregnancy specific beta-1-glycoprotein 6 M31125 1.30
 Consensus includes gb:AW974816 /FEA=EST /DB_XREF=gi:8166019 /DB_XREF=est:EST368021 /UG=Hs.283517 ESTs. Weakly similar to ALU1_HUMAN ALU SUBFAMI AW974816 1.30
 KIAA0988 protein AL136608 1.30
 tumor necrosis factor (ligand) superfamily, member 13 BF448647 1.30
 Consensus includes gb:AK025206.1 /DEF=Homo sapiens cDNA; FLJ21553 fls, clone COL06329 /FEA=mRNA /DB_XREF=gi:10437671 /UG=Hs.306788 Homo sapiens cDNA AK025206 1.30
 hypothetical protein FLJ10819 NM_018156 1.30
 Consensus includes gb:AL109708.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMAGE 254679 /FEA=mRNA /DB_XREF=gi:5689825 /UG=Hs.284310 AL109708 1.30
 KIAA0903 protein AB020710 1.30
 Consensus includes gb:AL137285.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D2416 (from clone DKFZp434D2416) /FEA=mRNA /DB_XREF=gi:5607740 /UG=Hs.30 AL137285 1.30
 chitinin (chitinase) 2 AK026415 1.30
 nitric oxide synthase 1 (neuronal) U31466 1.30
 zinc finger protein 20 (K0X13) AU154474 1.30
 tenascin XB X71923 1.30
 exotoxins (multiple)-like 3 BC006363 1.30
 Consensus includes gb:AL080134.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434G043 (from clone DKFZp434G043) /FEA=mRNA /DB_XREF=gi:5262575 /UG=Hs.1687 AL080134 1.30
 C-terminal binding protein 2 BC002486 1.30
 diubiquitin NM_006398 1.30
 EphB3 NM_004443 1.30
 solute carrier family 6 (neurotransmitter transporter, creatine), member 10 U41163 1.30
 ATP-binding cassette, sub-family G (WHITE), member 1 U34919 1.30
 folate receptor 2 (fetal) NM_000803 1.30
 HMT1 hnRNP methyltransferase-like 3 (S. cerevisiae) NM_019854 1.30
 splicing factor 30, survival of motor neuron-related BF224259 1.30
 ectodermal dysplasia 1, anhidrotic NM_001359 1.30
 pyruvate kinase, liver and RBC M15465 1.30
 estrogen receptor 1 AF238450 1.30
 formyl peptide receptor-like 1 M88107 1.30
 hypothetical protein FLJ10811 NM_018228 1.30
 calpastatin BG111635 1.30
 hypothetical protein FLJ13381 NM_025068 1.30
 KIAA0310 protein NM_025154 1.30
 zinc finger protein 232 NM_014519 1.30
 GAS2-related on chromosome 22 Y07846 1.30
 mucin 3A, intestinal AF007194 1.30
 Consensus includes gb:AK026465.1 /DEF=Homo sapiens cDNA; FLJ22812 fls, clone KIAA2955 /FEA=mRNA /DB_XREF=gi:10439335 /UG=Hs.306853 Homo sapiens cDNA AK026465 1.30
 hydroxysteroid (11-beta) dehydrogenase 2 NM_000196 1.30
 DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated) X98743 1.30
 catenin (cadherin-associated protein), alpha 2 M54886 1.30
 interferon, alpha 2 NM_005641 1.30
 thionexin-like 2 AV705253 1.30
 ankyrophil lateral sclerosis 2 (juvenile) chromosome region, candidate 3 NM_000696 1.30
 aldehyde dehydrogenase 9 family, member A1 NM_030930 1.30
 unc53 (C.elegans) homolog B W60953 1.30
 valosin-containing protein NM_001112 1.30
 adenosine deaminase, RNA-specific, B1 (homolog of rat RED1) NM_001112 1.30
 par-3 partitioning defective 3 homolog (C. elegans) AF197684 1.30
 chromosome 1 topen reading frame 9 NM_013279 1.30
 uretase homolog (Orosophila) AK000721 1.30
 RAB1, member RAS oncogene family AL530284 1.30
 thyroid stimulating hormone, beta NM_000549 1.30
 hypothetical protein MGC11271 NM_024323 1.30
 claudin 10 NM_006984 1.30
 translocase of inner mitochondrial membrane 17 homolog A (yeast) AK023063 1.30
 Consensus includes gb:AK027151.1 /DEF=Homo sapiens cDNA; FLJ23498 fls, clone LNG02683 /FEA=mRNA /DB_XREF=gi:10440206 /UG=Hs.306904 Homo sapiens cDNA AK027151 1.30
 hypothetical protein FLJ22390 NM_022746 1.30
 apolipoprotein B mRNA editing enzyme, catalytic polypeptide 1 NM_001644 1.30
 transforming growth factor, beta 3 J03241 1.30
 Consensus includes gb:AA587390 /FEA=EST /DB_XREF=est:mg82h01.s1 /CLONE=IMAGE:1090417 /UG=Hs.321110 ESTs. Highly similar to mucin 10 AA587390 1.30
 ret finger protein-like 3 antisense AL021837 1.30
 hypothetical protein FLJ14001 NM_024677 1.30
 B9 protein BC002944 1.30
 Consensus includes gb:AK028723.1 /DEF=Homo sapiens cDNA; FLJ23070 fls, clone LNG05829 /FEA=mRNA /DB_XREF=gi:10439841 /UG=Hs.305579 Homo sapiens cDNA AK028723 1.30

matrix metalloproteinase 26
EphA4
gdp junction protein, beta 3, 31KD (connexin 31)
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 7
KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3
PDZ and LIM domain 1 (elfin)
RGC32 protein
patently expressed 10
kinesin family member 1B
guanylate cyclase 1, soluble, beta 3
Alk-1 protein
myeloid cell leukemia sequence 1 (BCL2-related)
unc-5 homolog B (C. elegans)
gene for serine/threonine protein kinase
L-3-hydroxyacyl-Coenzyme A dehydrogenase, short chain
potassium voltage-gated channel, shaker-related subfamily, member 5
receptor (calcitonin) activity modifying protein 1
homo box D4
DKFZP364N1383 protein
cancer/testis antigen
mitogen-activated protein kinase kinase 6
integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)
hypothetical protein GL012
canine acyltransferase
transporter 2, ATP-binding cassette, sub-family B (MDR/TAP)
zinc finger protein 239
B-cell CLL/lymphoma 3
Consensus includes gb:BF222483 /FEA=EST /DB_XREF=gi:11129660 /DB_XREF=est:p53a01.x1 /CLONE=IMAGE:3649417 /UG=Hs.274454 ESTs, Weakly similar to 20X
Al157418
Consensus includes gb:AL117447.1 /DEF=Homo sapiens mRNA, cDNA DKFZp5641916 (from clone DKFZp5641916) /FEA=mRNA /DB_XREF=gi:4499947 /UG=Hs.3062
AL049218
CD59 antigen p18-20 (antigen identified by monoclonal antibodies 16.3A5, EJ16, EJ30, EJ32 and G344)
Consensus includes gb:AL049218.1 /DEF=Homo sapiens mRNA, cDNA DKFZp5641916 (from clone DKFZp5641916) /FEA=mRNA /DB_XREF=gi:4499947 /UG=Hs.3062
AL049218
interleukin 2 receptor, beta
glycerol kinase pseudogene 2
small nuclear ribonucleoprotein 70KD polypeptide (RNP antigen)
RNA-binding protein gene with multiple splicing
hypothetical protein FLJ22184
Consensus includes gb:AL117447.1 /DEF=Homo sapiens mRNA, cDNA DKFZp588A0617 (from clone DKFZp588A0617) /FEA=mRNA /DB_XREF=gi:5911895 /UG=Hs.304
AL117447
E7A-like factor 3 (ets domain transcription factor, epithelial-specific)
ets variant gene 1
hexophosphalidic acid acyltransferase-delta
NS1-associated protein 1
hypothetical protein FLJ14033 similar to hypoxia inducible factor 3, alpha subunit
H factor (complement)-like 3
sperm adhesion molecule 1 (PH-20 hyaluronidase, zona pellucida binding)
hydroxysteroid (11-beta) dehydrogenase 1
phosphodiesterase 8G, cGMP-specific, rod, gamma
potassium voltage-gated channel, KQT-like subfamily, member 2
protease, serine, 18 (thymus)
zinc finger protein ANC_2H01
T cell receptor alpha locus
HIC1-related gene on chromosome 22
SH2 domain-containing phosphatase E3 dunce homolog, Drosophila
histamine H4 receptor
tumor necrosis factor (ligand) superfamily, member 18
epididymal sperm binding protein 1
gb:U03891.2 /DEF=Homo sapiens phorbol 1 mRNA, complete cds. /FEA=mRNA PRODB=phorbol 1 /DB_XREF=gi:4895107 /UG=Hs.226307 phorbol (similar to apolipop
Consensus includes gb:BF434846 /FEA=EST /DB_XREF=gi:11447134 /DB_XREF=est:7074d07.x1 /CLONE=IMAGE:3641917 /UG=Hs.296757 Homo sapiens cDNA: FLJ2
Consensus includes gb:S78281.1 /DEF=pancreatic ribonuclease human, mRNA Recombinant Partial, 491 nt. /FEA=mRNA /GEN=pancreatic ribonuclease, HP-RNase /PRC
Consensus includes gb:AL109706.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 362430. /FEA=mRNA /DB_XREF=gi:5689823 /UG=Hs.32768
Consensus includes gb:AL442084.1 /DEF=Homo sapiens mRNA, cDNA DKFZp547G044 (from clone DKFZp547G044) /FEA=mRNA /DB_XREF=gi:10241763 /UG=Hs.308
AL442084
catechol-O-methyltransferase
RAB, member of RAS oncogene family-like 2A

Consensus includes gb:A1749193 /FEA=EST /DB_XREF=gi:127457 /DB_XREF=est:at0e04.x1 /CLONE=IMAGE:2374494 /UG=Hs.17639 Homo sapiens ubiquitin protein A1749193
 ubiquitin-activating enzyme E1C (UBA3) homolog, yeast) AL117566
 Consensus includes gb:BF766940 /FEA=EST /DB_XREF=gi:12101984 /DB_XREF=est:602258153F1 /CLONE=IMAGE:4341588 /UG=Hs.294100 ESTs BF766940
 carboxypeptidase A1 (pancreatic) NM_001868
 glycine-N-acyltransferase AW024233
 survival motor neuron pseudogene U05682
 erythrocyte membrane protein band 4.2 U81640
 ATPase, Ca++ transporting, type 2C, member 1 M30646
 hypothetical protein FLJ22087 AF225881
 cyclin D1 (PRAD1; parathyroid adenomatosis 1) A4555088
 chondroilin sulfate proteoglycan 2 (versican) M73554
 Consensus includes gb:AK026793.1 /DEF=Homo sapiens cDNA: FLJ23140 fs, clone LNG09065, /FEA=mRNA /DB_XREF=gi:10439729 /UG=Hs.287728 Homo sapiens cDNA: AK026793
 Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse) M92424
 potassium voltage-gated channel, shaker-related subfamily, beta member 2 AL520102
 Consensus includes gb:A1912086 /FEA=EST /DB_XREF=est:wd8408.x1 /CLONE=IMAGE:233831 /UG=Hs.194750 Human DNA sequence from cA1912086
 syndecan 2 (heparan sulfate proteoglycan 1, cell surface-associated, fibroglycan) BE048514
 Consensus includes gb:AW301235 /FEA=EST /DB_XREF=gi:6710912 /DB_XREF=est:xs69c06.x1 /CLONE=IMAGE:2774890 /UG=Hs.66185 Homo Sapiens mRNA, partial AW301235
 golgi associated, gamma adaptin ear containing, ARF binding protein 2 AW088154
 CD86 antigen (CD28 antigen ligand 2, B7-2 antigen) L25259
 ATP synthase, H+ transporting, mitochondrial F0 complex, subunit 9 AA917672
 chromosome 11 open reading frame 14 NM_020645
 karyopherin alpha 8 (importin alpha 7) BG403834
 Cluster incl. A1738452; w32a07.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2391924 /clone_end=3 /gb=A1738452 /gi=5100433 /ug=Hs.203286 /len=533 A1738452
 leucine rich repeat (in FLII) interacting protein 1 AF130054
 hydroxyteroid (17-beta) dehydrogenase 7 NM_016371
 formyl peptide receptor 1 M60626
 protein phosphatase 1B (formery 2C), magnesium-dependent, beta isoform AJ271832
 glycoprotein M8A BF939489
 Consensus includes gb:A207792 /FEA=EST /DB_XREF=gi:3769734 /DB_XREF=est:ao89403.x1 /CLONE=IMAGE:1953029 /UG=Hs.306019 ESTs, Weakly similar to ALU7 AC06271
 golgi phosphoprotein 1 NM_022735
 fibronectin II NM_012189
 mammoside, alpha, class 2C, member 1 NM_006715
 ets variant gene 6 (TEL oncogene) AF041811
 v-myc myelocytomatosis viral oncogene homolog 1, lung carcinoma derived (avian) M19720
 KIA0673 protein A117405
 Consensus includes gb:AK024457.1 /DEF=Homo sapiens mRNA for FLJ00049 protein, partial cds, /FEA=mRNA /GEN=KRTAP2.1A /PROD=keratin associated protein K NM_030977
 Consensus includes gb:AUI47820 /FEA=EST /DB_XREF=gi:11009141 /DB_XREF=est:AUI47820 /CLONE=MAMMA1001206 /UG=Hs.288490 Homo sapiens cDNA FLJ12 AUI47820
 double-stranded RNA specific adenosine deaminase NM_018702
 hypothetical protein FLJ13031 NM_024688
 Consensus includes gb:A039469 /FEA=EST /DB_XREF=gi:3278663 /DB_XREF=est:ox41a08.s1 /CLONE=IMAGE:1658870 /UG=Hs.27362 ESTs A039469
 aquaporin 8 NM_001169
 Consensus includes gb:AK024995.1 /DEF=Homo sapiens cDNA: FLJ21342 fs, clone COL02873, /FEA=mRNA /DB_XREF=gi:10437426 /UG=Hs.308739 Homo sapiens cDNA: AK024995
 a disintegrin and metalloproteinase domain 7 AF215624
 phospholipase D1, phosphatidylcholine-specific U38545
 hypothetical protein DKFZ043AC0923 NM_017598
 Consensus includes gb:AL035587 /DEF=Human DNA sequence from clone RP3-475N16 on chromosome 6p12.3-21.2. Contains the genes for CTGA4, pre-T cell receptor, BF115424
 progesterin induced protein AF119817
 discs, large (Drosophila) homolog-associated protein 2 NM_024689
 hypothetical protein FLJ14103 AF052146
 Consensus includes gb:AF052146.1 /DEF=Homo sapiens clone 24653 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3380455 /UG=Hs.155085 Homo sapiens clone 24653 AF052146
 KIA0523 protein BF115148
 disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila) NM_001343
 flavin containing monooxygenase 5 NM_001461
 hepatocyte nuclear factor 3, beta AB028021
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1 NM_003069
 tripartite motif-containing 10 AF220122
 cleavage and polyadenylation specific factor 1, 160KD subunit BE676642
 hypothetical protein DKFZp762B226 AK001821
 galactosylceramidase (Krabbe disease) NM_000153
 chromosome 1 open reading frame 14 NM_030933

protein tyrosine phosphatase, receptor type, F	1,29	ALU145351
interleukin 9 receptor	1,29	L39064
death-associated protein kinase 1	1,29	NM_004938
hypothetical protein PRO1787	1,29	NM_018606
signal transducer and activator of transcription 1, p1kD	1,29	M87835
olfactory receptor, family 1, subfamily E, member 2	1,29	NM_003554
surfactant protein A binding protein	1,29	NM_005129
collin 4A	1,29	NM_003569
KIAA0918 protein	1,29	AW449813
Bacteriophage IREF-X03453 DEF= Bacteriophage P1 are recombinase protein corresponding to nucleotides 581-1001 of X03453 /LEN= 1058 (-5 and -3 represent transcrit	1,29	X03453
Consensus includes gb:AF035314.1 /DEF= Homo sapiens clone 23651 mRNA sequence. /FEA= mRNA /DB_ XREF= gi:2661076 /UG= Hs.134528 Homo sapiens clone 23651	1,29	AF035314
claudin 18	1,29	NM_016369
DKFZP434C212 protein	1,29	AK023841
ephrin-A2	1,29	NM_001405
chondroitin sulfate proteoglycan BEHAB/brevican	1,29	A205180
KIAA0231 protein	1,29	R41496
aldo-keto reductase family 1, member B10 (aldose reductase)	1,29	NM_020299
dynen, axonemal, heavy polypeptide 17	1,29	AL122077
Consensus includes gb:AF035382 /FEA= EST /DB_ XREF= gi:989223 /DB_ XREF= est:yd19506.s1 /CLONE= IMAGE:274115 /UG= Hs.124984 ESTs. Moderately similar to ALU7_	1,29	H49392
collagen, type XIII, alpha 1	1,29	M33653
Ras suppressor protein 1	1,29	NM_012425
frizzled homolog 7 (Drosophila)	1,29	NM_003507
secreted protein, acidic, cysteine-rich (osteonectin)	1,29	NM_003118
kinesin family member 3B	1,29	NM_004788
sarcosine dehydrogenase	1,29	NM_007101
hypothetical protein FLJ21887	1,29	NM_024659
anaplastic lymphoma kinase (Kt-1)	1,29	NM_004304
adaptor protein containing pH domain, PTB domain and leucine zipper motif	1,29	NM_012096
Consensus includes gb:BF431618 /FEA= EST /DB_ XREF= gi:11443732 /DB_ XREF= est:7016603.x1 /CLONE= IMAGE:3574349 /UG= Hs.12342 Homo sapiens clone 24538_	1,29	BF431618
glutamate receptor, ionotropic, N-methyl D-aspartate 2C	1,29	L76224
Consensus includes gb:AL050335 /DEF= Human DNA sequence from clone RP1-190J20 on chromosome 6p22.3-24.1. Contains ESTs, STSs and GSSs. Contains a putative	1,29	AL050335
Cytochrome P450, subfamily 11E (ethanol-inducible)	1,29	AF182276
adenosine A3 receptor	1,29	NM_000677
hypothetical protein FLJ20234	1,29	BC000785
alpha-2-glycoprotein 1, zinc	1,29	AC004522
Consensus includes gb:AF070577.1 /DEF= Homo sapiens clone 24461 mRNA sequence. /FEA= mRNA /DB_ XREF= gi:3387949 /UG= Hs.4817 Homo sapiens clone 24461	1,29	AF070577
1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)	1,29	AF007145
eukaryotic translation initiation factor 2, subunit 1 (alpha, 35kD)	1,29	A4577688
glial axonal neuropathy (gigaxonin)	1,29	NM_022041
CD209 antigen-like	1,29	AF245219
Fc fragment of IgE, low affinity II, receptor for (CD23A)	1,29	NM_002002
hypothetical protein FLJ21834	1,29	NM_022087
Consensus includes gb:AL034398 /DEF= Human DNA sequence from clone 1158B12 on chromosome Xp11.21-11.4 Contains the ZXDA gene for X-linked duplicated Zinc	1,29	AL034398
transforming growth factor, beta 2	1,29	BF061658
polycystic kidney disease 1 (autosomal dominant)	1,29	L39891
chromosome 1 open reading frame 6	1,29	NM_004872
putative integral membrane transporter	1,29	T15777
Consensus includes gb:AK022362.1 /DEF= Homo sapiens cDNA FLJ12300 fts, clone MAMMA1001854. /FEA= mRNA /DB_ XREF= gi:10433744 /UG= Hs.287510 Homo sapie	1,29	AK022362
tumor-associated calcium signal transducer 1	1,29	NM_002354
Consensus includes gb:AK024871.1 /DEF= Homo sapiens cDNA: FLJ121218 fts, clone COL00537. /FEA= mRNA /DB_ XREF= gi:10437282 /UG= Hs.306712 Homo sapiens cD	1,29	AK024871
TUJ3A protein	1,29	NM_007177
H1 histone family, member 5	1,29	NM_005322
fatty-acyl-Coenzyme A ligase, long-chain 5	1,29	NM_018234
histamine N-methyltransferase	1,29	N40285
synuclein, gamma (breast cancer-specific protein 1)	1,29	AF010128
claudin 6	1,29	NM_021195
KIAA0877 protein	1,29	AB020684
putative DNA/chromatin binding motif	1,29	NM_006818
Consensus includes gb:AW241752 /FEA= EST /DB_ XREF= gi:6575506 /DB_ XREF= est:xn74404.x1 /CLONE= IMAGE:2700223 /UG= Hs.160728 Homo sapiens cDNA FLJ116	1,29	AW241752
gap junction protein, beta 5 (connexin 31.1)	1,29	NM_005288
matrix metalloproteinase 14 (membrane-inserted)	1,29	Z48481
hypothetical protein FLJ22021	1,29	NM_024535
tracr	1,29	AF106861

protein phosphatase 1, regulatory (inhibitor) subunit 3C
interferon regulatory factor 1
Src-like-adapter
DnaJ (Hsp40) homolog, subfamily C, member 4
DKFZP434D156 protein
cyclin D2
Consensus includes gb:A822538 /FEA=EST /DB_XREF=gi:5658502 /DB_XREF=est:wm89c11.x1 /CLONE=IMAGE:2443124 /UG=Hs.152860 Homo sapiens mRNA; cDNA A822538
dentatorubral-pallidolysian atrophy (drip) (atrophin-1)
WW domain-containing oxidoreductase
protocadherin 16
TERA protein
hypothetical protein MGC4090
L1 cell adhesion molecule (hydrocephalus, stenosis of aqueduct of Sylvius 1, MASA (mental retardation, aphasia, shuffling gait and adducted thumbs) syndrome, spastic pa
hypothetical protein MGC4294
nuclear RNA export factor 3
hypothetical protein FLJ21044 similar to Rbg1
immunoglobulin superfamily containing leucine-rich repeat
adrenergic, alpha-1A-, receptor
carbohydrate (N-acetylglucosamine 6-O-) sulfotransferase 4
t-complex 10 (a murine top homolog)
hypothetical protein FLJ20126
LIM domain kinase 2
Consensus includes gb:BF511878 /FEA=EST /DB_XREF=gi:11594976 /DB_XREF=est:UH-BH-aom-c-02-0-U1.s1 /CLONE=IMAGE:3085418 /UG=Hs.128311 ESTs
four and a half LIM domains 3
Consensus includes gb:AL080082.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G1182 (from clone DKFZp564G1182) /FEA=mRNA /DB_XREF=gi:5262495 /UG=Hs.13
transformer-2 alpha (htra-2 alpha)
endothelin converting enzyme-like 1
transient receptor potential channel 4
hypothetical protein FLJ11526
Kruppel-like factor 3 (basic)
protein 2, vitamin K-dependent plasma glycoprotein
carbohydrate kinase-like
Consensus includes gb:AV651668 /FEA=EST /DB_XREF=gi:9872682 /DB_XREF=est:AV651668 /CLONE=GLCSC04 /UG=Hs.282480 ESTs
lymphocyte activation-associated protein
oxidative 3 alpha hydroxysteroid dehydrogenase; retinol dehydrogenase; 3-hydroxysteroid epimerase
PAB28, member RAS oncogene family
hypothetical protein FLJ20515
MAP1/microtubule affinity-regulating kinase 3
hydroxysteroid (17-beta) dehydrogenase 3
triophosphate isomerase 1
tropomyosin 1 (alpha)
zinc finger protein 258
hypothetical protein FLJ12704
Fc fragment of IgG, high affinity (a, receptor for (CD64)
growth associated protein 43
interleukin 1 receptor accessory protein-like 1
Consensus includes gb:BF223401 /FEA=EST /DB_XREF=gi:11130578 /DB_XREF=est:7q87n04.x1 /CLONE=IMAGE:3705462 /UG=Hs.112158 Homo sapiens PAC clone F BF223401
calctinin, gene-related peptide-receptor component protein
KIAA0582 protein
pre-alpha (globulin) inhibitor, H3 polypeptide
CCR4-NOT transcription complex, subunit 7
alcohol dehydrogenase 7 (class IV), mu or sigma polypeptide
protein tyrosine phosphatase type IVA, member 1
inositol 1,3,4-trisphosphate 5b kinase
complement component (3b/4b) receptor 1, including Knops blood group system
cadherin 8, type 2
Consensus includes gb:AL080104.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566D153 (from clone DKFZp566D153) /FEA=mRNA /DB_XREF=gi:5262529 /UG=Hs.3063
allograft inflammatory factor 1
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson
hypothetical protein FLJ20220
Consensus includes gb:AJ005205 /DEF=Homo sapiens 5HT3 gene for serotonin 3 receptor /FEA=mRNA_2 /DB_XREF=gi:7019744 /UG=Hs.2142 5-hydroxytryptamine (ser A005205
Consensus includes gb:AK000244.1 /DEF=Homo sapiens cDNA FLJ20237 fls, clone COLF-5879, highly similar to AB023230 Homo sapiens mRNA for KIAA1013 protein. fl AK000244

plateletin 4	AA194254	1.28
hyaluronan-mediated motility receptor (RHAMM)	NM_012485	1.28
gamma-aminobutyric acid (GABA) A receptor, alpha 1	NM_000806	1.28
syntaxin 11	AF071504	1.28
crystallin, alpha B	AF007162	1.28
Consensus includes gb:AW838210 /FEA=EST /DB_XREF=gi:7930184 /DB_XREF=est:RC3-LT0023-200100-012-e09 /UG=Hs.278648 Homo sapiens cDNA FLJ14085 fs. (AW838210)	1.28	
internally expressed 3	AI950273	1.28
ribonuclease, RNase A family, k6	NM_005615	1.28
prostaglandin O2 synthase (21kD, brain)	NM_000954	1.28
Consensus includes gb:AL049285.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564M193 (from clone DKFZp564M193) /FEA=mRNA /DB_XREF=gi:4500045 /UG=Hs.302C	1.28	
defensin, beta 1	U73945	1.28
superoxide dismutase 3, extracellular	NM_003102	1.28
cytochrome P450, subfamily 11B (phenobarbital-inducible), polypeptide 6	NM_000767	1.28
AF15q14 protein	NM_020380	1.28
receptor tyrosine kinase-like orphan receptor 2	NM_004560	1.28
aquaporin 1 (channel-forming integral protein, 28kD)	NM_000385	1.28
Consensus includes gb:H15535 /FEA=EST /DB_XREF=gi:880355 /DB_XREF=est:ym27c01.s1 /CLONE=IMAGE:49385 /UG=Hs.52792 Homo sapiens mRNA: cDNA DKFZ: H15535	1.28	
homeo box C8	NM_004503	1.28
type 1 transmembrane protein Fn14	NM_016639	1.28
glutathione S-transferase M2 (muscle)	NM_000848	1.28
butyrophilin, subfamily 3, member A3	NM_006894	1.28
NADPH oxidase 1	AF168327	1.28
group III secreted phospholipase A2	NM_015715	1.28
synaptotagmin 1	NM_004711	1.28
tumor endothelial marker 8	NM_018153	1.28
KIAA1107 protein	AB029030	1.28
cytochrome c oxidase subunit IV isoform 1	AW337510	1.28
spindlin-like	NM_019003	1.28
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, 812)	NM_02491	1.28
KIAA1048 protein	BC002695	1.28
trans-phenyltransferase	NM_014317	1.28
ret finger protein-like 1	NM_021026	1.28
splicing factor, arginine/serine-rich 6	NM_006275	1.28
translocation protein 1	BE866511	1.28
CD44 antigen (homing function and Indian blood group system)	BC004372	1.28
procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase) 2	NM_000935	1.28
chloride channel, calcium activated, family member 1	AF127036	1.28
D-aspartate oxidase	NM_003649	1.28
paraoxonase 1	NM_000446	1.28
matrix metalloproteinase 27	NM_022122	1.28
glycoprotein Ib (platelet), alpha polypeptide	NM_000173	1.28
hypothetical protein FLJ20254	NM_017727	1.28
like-glycosyltransferase	NM_004737	1.28
Consensus includes gb:AL157490.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434F1011 (from clone DKFZp434F1011) /FEA=mRNA /DB_XREF=gi:7018535 /UG=Hs.27: M33318	1.28	
cytochrome P450, subfamily 11A (phenobarbital-inducible), polypeptide 6	1.28	
Consensus includes gb:AL359557.1 /DEF=Homo sapiens mRNA: cDNA DKFZp762O1415 (from clone DKFZp762O1415) /FEA=mRNA /DB_XREF=gi:8655612 /UG=Hs.30	1.28	
fibroblast growth factor receptor 3 (achondroplasia, thanatophoric dwarfism)	NM_000142	1.28
KIAA0485	AI073551	1.28
cell division cycle 2-like 5 (cholinesterase-related cell division controller)	BC001274	1.28
Consensus includes gb:AU145003 /FEA=EST /DB_XREF=gi:11006524 /DB_XREF=est:AU145003 /CLONE=HEMBA1003598 /UG=Hs.287428 Homo sapiens cDNA FLJ111	1.28	
gb:NM_020484.1 /DEF=Homo sapiens RAGE binding protein (AF011757), mRNA, /FEA=mRNA /GEN=AF011757 /PROD=RAGE binding protein /DB_XREF=gi:10048411 / NM_020484	1.28	
erythrocyte membrane protein band 4.9 (dema1n)	NM_001978	1.28
phosphodiesterase 8A	W73272	1.28
interleukin 1 receptor antagonist	U65560	1.28
ATPase, Ca++ transporting, cardiac muscle, slow twitch 2	AA805753	1.28
5-hydroxytryptamine (serotonin) receptor 1B	M81590	1.28
sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3C	AI962897	1.28
transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)	AI871234	1.28
Consensus includes gb:AF043583.1 /DEF=Homo sapiens clone ASMmeg1-63 immunoglobulin lambda chain VJ region, (JGL) mRNA, partial cds. /FEA=mRNA /GEN=JGL /AF043583	1.28	
golgi associated, gamma adaptin ear containing, ARF binding protein 3	AF218138	1.28
Eeg-related gene member 2	NM_030779	1.28
adenyosuccinate lyase	AL022236	1.28
hypothetical protein FLJ10916	NM_018271	1.28

hypothetical protein MGC5457
 afactomedin 1
 solute carrier family 31 (copper transporters), member 2
 WD repeat domain 12
 LIM domain protein
 solute carrier family 17 (anion/sugar transporter), member 5
 hypothetical protein from clone 24796
 disrupted in schizophrenia 1
 putative DNA/chromatin binding motif
 FN5 protein
 wingless-type MMTV integration site family, member 6
 DnaJ (Hsp40) homolog, subfamily B, member 5
 Mdm4, transformed 373 cell double minute 4, p53 binding protein (mouse)
 cathepsin G
 KIAA1117 protein
 fibron (collagen/fibrinogen domain-containing) 1
 hypothetical protein MGC4683
 runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene)
 phospholipase C, beta 4
 mitogen-activated protein kinase kinase 4
 NAD kinase
 forthead box F1
 acetyl LDL receptor; SREC=scavenger receptor expressed by endothelial cells
 melanoma antigen, family C, 1
 tryptophan 2,3-dioxygenase
 hypothetical protein FLJ12988
 inositol 1,4,5-triphosphate receptor, type 2
 solute carrier family 25 (mitochondrial carrier; peroxisomal membrane protein, 34kD), member 17
 heme oxygenase (decycling) 1
 Cockayne syndrome 1 (classical)
 deleted in azoospermia 3
 homolog of rat orphan transporter v7-3
 TNFRSF1A-associated via death domain
 dermatopontin
 small muscle protein, X-linked
 inhibitor of growth family, member 1
 olfactory receptor, family 1, subfamily D, member 2
 DKFZP434A043 protein
 cytochrome b-245, beta polypeptide (chronic granulomatous disease)
 angiotensin-like 4
 hypothetical protein FLJ22202
 PK506 binding protein 9 (63 kD)
 Consensus includes gb:AL049260.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564E233 (from clone DKFZp564E233). /FEA=mRNA /DB_XREF=gi:4500007 /UG=Hs.3020
 eukaryotic translation initiation factor 4 gamma, 3
 dynein, axonemal, light intermediate polypeptide
 SRY (sex determining region Y)-box 18
 3-hydroxybutyrate dehydrogenase (heart, mitochondrial)
 serine/threonine kinase 38
 KIAA0328 protein
 uracil-DNA glycosylase
 Consensus includes gb:U95737 /DEF=Human Chromosome 16 BAC clone CIT987SK-A-388D4 /FEA=mRNA /DB_XREF=gi:3417290 /UG=Hs.177961 Human Chromosome
 differentiation-related protein dif13
 transmembrane protease, serine 5 (spinesin)
 growth factor receptor-bound protein 14
 proline and glutamic acid rich nuclear protein
 hypothetical protein FLJ13081
 isocitrate dehydrogenase 2 (NADP+), mitochondrial
 phosphoinositid 3-phosphate-binding protein-2
 tachykinin receptor 3
 C-reactive protein, pentraxin-related
 membrane protein, palmitoylated 2 (MAGUK p55 subfamily member 2)
 transmembrane 4 superfamily member 1
 myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 2

hypothetical protein FLJ14126
 Consensus includes gb:AK021576.1 /DEF=Homo sapiens cDNA FLJ11514 f1s, clone HEMBA1002229, /FEA=mRNA /DB_XREF=gi:10432783 /UG=Hs.306602 Homo sapiens
 polydomyate binding protein-interacting protein 1
 ELAV (embryonic lethal, abnormal vision, Drosophila)-like 2 (Hu antigen B)
 Consensus includes gb:AA131324 /FEA=EST /DB_XREF=gi:1693011 /DB_XREF=est:237b04.s1 /CLONE=IMAGE:503593 /UG=Hs.28264 Homo sapiens mRNA; cDNA DK AA131324
 ras homolog gene family, member E
 DNA segment on chromosome X and Y (unique) 155 expressed sequence
 CCAAT/enhancer binding protein (C/EBP) delta
 Consensus includes gb:AK025194.1 /DEF=Homo sapiens cDNA: FLJ21541 f1s, clone COL06166, /FEA=mRNA /DB_XREF=gi:10437657 /UG=Hs.306784 Homo sapiens cDNA
 DKFZP566A0522 protein
 hypothetical protein FLJ23074
 myosin, heavy polypeptide 11, smooth muscle
 PAN2 protein
 astrotactin 2
 dystroglycan 1 (dystrophin-associated glycoprotein 1)
 general transcription factor IIA, 1 (37KD and 19KD subunits)
 potassium voltage-gated channel, Shab-related subfamily, member 1
 glycoprotein M6B
 Consensus includes gb:AL049285.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564F053 (from clone DKFZp564F053), /FEA=mRNA /DB_XREF=gi:4500013 /UG=Hs.71981 AL049285
 matrix metalloproteinase 12 (macrophage elastase)
 TGFB inducible early growth response
 Consensus includes gb:AU148255 /FEA=EST /DB_XREF=gi:1009776 /DB_XREF=est:AU148255 /CLONE=MAMMA1003031 /UG=Hs.287627 Homo sapiens cDNA FLJ14
 hypothetical protein FLJ10312
 zinc finger protein 165 (LIM domain)
 beta-galactose-3-O-sulfotransferase, 4
 PRO233 protein
 tissue inhibitor of metalloproteinase 1 (erythroid potentiating activity, collagenase inhibitor)
 paired immunoglobulin-like receptor alpha
 atrophin-1 interacting protein 1; activin receptor interacting protein 1
 hypothetical protein
 Consensus includes gb:X63966.1 /DEF=H sapiens synthetic gene for platelet-derived growth factor-BB, /FEA=mRNA (PROD=platelet-derived growth factor-BB /DB_XREF=X63966
 go:U41742.1 /DEF=Human nucleophosmin-retinoid acid receptor alpha fusion protein NPM-RAR long form mRNA, complete cds, /FEA=mRNA (PROD=nucleophosmin-reti
 hypothetical protein FLJ20718
 KIAA1466 protein
 oxysterol-binding protein-related protein 1
 adaptor-related protein complex 1, gamma 2 subunit
 myotubularin related protein 3
 hypothetical protein MGC3020
 ubiquitin-conjugating enzyme E2L 6
 Consensus includes gb:AL137475.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434F0723 (from clone DKFZp434F0723), /FEA=mRNA /DB_XREF=gi:6808086 /UG=Hs.27
 angiotensin-like factor
 ret finger protein-like 3 antisense
 Misshapen/NIK-related kinase
 transcription termination factor, RNA polymerase II
 sialic acid binding Ig-like lectin 7
 ankyrin 1, erythrocytic
 zinc finger protein 133 (clone pHZ-13)
 hypothetical protein LOC37187
 hypothetical protein FLJ11506
 interferon gamma receptor 1
 serologically defined colon cancer antigen 1
 transcription factor 7 (T-cell specific, HMG-box)
 hypothetical protein MGC4293
 hypothetical protein FLJ20896
 KIAA0404 protein
 ferritin, light polypeptide
 thrombomodulin
 microsomal glutathione S-transferase 3
 X11L-binding protein 51
 hypothetical protein FLJ22002
 cytoskeleton associated protein 2
 endothelin converting enzyme 2
 Consensus includes gb:AU145806 /FEA=EST /DB_XREF=gi:1007327 /DB_XREF=est:AU145806 /CLONE=HEMBA1005894 /UG=Hs.286664 Homo sapiens cDNA FLJ11; AU145806

hypothetical protein FLJ20428	NM_017817	1.28
E74-like factor 5 (ets domain transcription factor)	AF115403	1.28
matrix metalloproteinase 14 (membrane-inserted)	NM_004995	1.28
hypothetical protein	AV700415	1.28
protein tyrosine phosphatase, receptor type, R	NM_002849	1.28
hypothetical protein MGC10902	NM_030570	1.28
homeo box A10	BF792917	1.28
KIAA0261 protein	D87450	1.28
tripartite motif-containing 3	NM_006458	1.28
ubiquitously transcribed tetratricopeptide repeat gene, X chromosome	AI140752	1.28
Consensus includes gb:AW056600 /FEA=EST /DB_XREF=gi:5934239 /DB_XREF=est:wx23h08.x1 /CLONE=IMAGE:2544543 /UG=Hs.18593 Homo sapiens cDNA: FLJ21	AW056600	1.28
peroxisome biogenesis factor 10	AB013818	1.28
SA (rat hypertension-associated) homolog	NM_005622	1.28
MAD, mothers against decapentaplegic homolog 2 (Drosophila)	AW151617	1.28
KIAA0934 protein	NZ2859	1.28
Consensus includes gb:AL122122.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434L098 (from clone DKFZp434L098). /FEA=mRNA /DB_XREF=gi:6102949 /UG=Hs.2745	AL122122	1.28
calpain 9 (pCL-4)	NM_008615	1.28
asporin (LRR class 1)	NM_017680	1.28
defensin, alpha 6, Paneth cell-specific	NM_001926	1.28
PC326 protein	NM_018442	1.27
nucleotide binding protein 1 (MinD homolog; E. coli)	NM_002484	1.27
zinc finger protein 14 (KOF 6)	NM_021030	1.27
transporter 1, ATP-binding cassette, sub-family B (MDR/TP)	NM_000593	1.27
Consensus includes gb:AL050032.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566F1224 (from clone DKFZp566F1224). /FEA=mRNA /DB_XREF=gi:4884272 /UG=Hs.30	AL050032	1.27
hypothetical protein FLJ20047	NM_017639	1.27
protease inhibitor 15	NM_015886	1.27
ryanodine receptor 3	NM_001036	1.27
retinal short-chain dehydrogenase/reductase retSDR2	NM_016245	1.27
LIM domain only 4	SC003600	1.27
Consensus includes gb:A1813483 /FEA=EST /DB_XREF=est:ty37e12.x1 /CLONE=IMAGE:2281284 /UG=Hs.324504 Homo sapiens mRNA: cDNA	AI813483	1.27
protein tyrosine phosphatase, receptor type, J	AI800462	1.27
KIAA0356 gene product	AB002354	1.27
transcription factor 7-like 2 (T-cell specific, HMG-box)	AI375916	1.27
T-cell acute lymphocytic leukemia 1	NM_003189	1.27
Consensus includes gb:AF218021.1 /DEF=Homo sapiens clone PP552 unknown mRNA. /FEA=mRNA /PROD=unknown /DB_XREF=gi:10441971 /UG=Hs.302127 Homo	AF218021	1.27
Wilms tumor 1	NM_024426	1.27
RAB2, member RAS oncogene family	AI743756	1.27
Consensus includes gb:N71739 /FEA=EST /DB_XREF=gi:1228451 /DB_XREF=est:tyw52h11.1 /CLONE=IMAGE:255909 /UG=Hs.237554 Homo sapiens cDNA FLJ14308	N71739	1.27
hypothetical protein FLJ22938	NM_024676	1.27
KRAB box containing C2H2 type zinc finger bA526D8.4	BE567977	1.27
FERM, RhoGEF (ARHGGEF) and pleckstrin domain protein 1 (chondrocyte-derived)	NM_005766	1.27
zinc finger protein 93 homolog (mouse)	NM_004234	1.27
Consensus includes gb:AL050032.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566F1224 (from clone DKFZp566F1224). /FEA=mRNA /DB_XREF=gi:4884272 /UG=Hs.30	AL050032	1.27
major histocompatibility complex, class II, DO alpha	M29335	1.27
G-2 and S-phase expressed 1	AI340239	1.27
DKFZP564G0222 protein	NM_014044	1.27
mitogen-activated protein kinase kinase 5	ZZ5426	1.27
gb:BC001957.1 /DEF=Homo sapiens, Similar to KIAA0144 gene product, clone MGC-761, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0144 gene product /	BC001957	1.27
Consensus includes gb:AK024315.1 /DEF=Homo sapiens cDNA FLJ14253 flt, clone OVARC1001376. /FEA=mRNA /DB_XREF=gi:10436666 /UG=Hs.300694 Homo sapie	AK024315	1.27
Consensus includes gb:AL390157.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D179 (from clone DKFZp434D179). /FEA=mRNA /DB_XREF=gi:9368847 /UG=Hs.3065	AL390157	1.27
MCF 2 cell line derived transforming sequence	X13230	1.27
Consensus includes gb:AK021614.1 /DEF=Homo sapiens cDNA FLJ11552 flt, clone HEMBA1003021. /FEA=mRNA /DB_XREF=gi:10432830 /UG=Hs.306606 Homo sapie	AK021614	1.27
collagen, type V, alpha 1	N30339	1.27
adaptor-related protein complex 1, gamma 1 subunit	NM_001128	1.27
tankyrase, TRF-1-interacting ankyrin-related ADP-ribose polymerase	AF082559	1.27
KIAA0316 gene product	AW297087	1.27
vasodilator-stimulated phosphoprotein	NM_003370	1.27
protein kinase (cAMP-dependent, catalytic) inhibitor alpha	NM_006823	1.27
hypothetical protein	BC005004	1.27
thymidine kinase 2, mitochondrial	BE895437	1.27
butyrophilin-like 3	AK025287	1.27
matrix metalloproteinase 8 (neutrophil collagenase)	NM_002424	1.27
BA11-associated protein 2	NM_017451	1.27

transcription factor 3 (E2A immunoglobulin enhancer binding factors E12/E47)
endothelin 3
KIAA0826 protein
TYRO3 protein tyrosine kinase
Consensus includes gb:AL117659.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586i0624 (from clone DKFZp586i0624) /FEA=mRNA /DB_XREF=gi:5912252 /UG=Hs.3086; AL117659
calcium channel, voltage-dependent, alpha 11 subunit
cholinergic receptor, nicotinic, alpha polypeptide 1 (muscle)
tumor necrosis factor (ligand) superfamily, member 4 (tax-transcriptionally activated glycoprotein 1, 34kD)
Consensus includes gb:AK000995.1 /DEF=Homo sapiens cDNA FLJ10133 fs, clone HEMBA1003087 /FEA=mRNA /DB_XREF=gi:7021998 /UG=Hs.272209 Homo sapiens
Consensus includes gb:AI191118 /FEA=EST /DB_XREF=gi:3742327 /DB_XREF=est:q16c05.x1 /CLONE=IMAGE:1739336 /UG=Hs.222015 ESTs
stromatin (EBP72)-like 1
Consensus includes gb:AL031686 /DEF=Human DNA sequence from clone 981L23 on chromosome 20q12.1-13.2. Contains a Krueppel type zinc-finger protein pseudogen
ribonuclease, RNase A family, 4
cardiotrophin-like cytokine: neurotrophin-1/B-cell stimulating factor-3
hypothetical protein FLJ10598
leukocyte membrane antigen
LIM domain binding 3
KIAA0061 protein
methylmalonyl Coenzyme A mutase
Consensus includes gb:R25849 /FEA=EST /DB_XREF=gi:781984 /DB_XREF=est:Y42a11.r1 /CLONE=IMAGE:132380 /UG=Hs.217305 ESTs
TBP-like 1
KIAA0671 gene product
KIAA0878 protein
Consensus includes gb:AF238870.1 /DEF=Homo sapiens clone GLSH-3 similar to gliadin mRNA sequence. /FEA=mRNA /DB_XREF=gi:8272577 /UG=Hs.275706 Homo sapiens
myelin-associated oligodendrocyte basic protein
FGFR1 oncogene partner
protocadherin gamma subfamily A, 1
Consensus includes gb:AX023116.1 /DEF=Homo sapiens cDNA FLJ13054 fs, clone NT2RP3001527, highly similar to Human Sp140 protein (Sp140) mRNA. /FEA=mRNA /DB_XREF=gi:8272577 /UG=Hs.275706 Homo sapiens
hemochromatosis
light junction protein 1 (zone occludens 1)
connexin-36
Consensus includes gb:AL335942.1 /DEF=Homo sapiens mRNA; cDNA DKFZp76L23121 (from clone DKFZp76L23121) /FEA=mRNA /DB_XREF=gi:7669981 /UG=Hs.275706 Homo sapiens
lipocalin 2 (oncogene 24p3)
hypothetical protein FLJ11467
Consensus includes gb:AF107846 /DEF=Homo sapiens neuroendocrine-specific Golgi protein p55 (XLalpha) gene, exon XL2 and complete cds /FEA=mRNA /DB_XREF=gi:7669981 /UG=Hs.275706 Homo sapiens
RAB interacting factor
RAB binding protein 1
plexin A2
interleukin-1 homolog 1
ectonucleoside triphosphate diphosphohydrolase 2
estrogen-related receptor gamma
Consensus includes gb:AI375002 /FEA=EST /DB_XREF=gi:4174992 /DB_XREF=est:ts4b11.x1 /CLONE=IMAGE:2047869 /UG=Hs.272014 ESTs. Weakly similar to ALU3 /AI375002
collagen, type VI, alpha 1
lymphocyte antigen 95 (activating NK-receptor, NK-p44)
KIAA0984 protein
syntaxin 6
TAF7 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 55 kD
KIAA0276 protein
serine (or cysteine) proteinase inhibitor, clade E (nexin, plasminogen activator inhibitor type 1), member 1
X03453 Bacteriophage P1 cre recombinase protein (-5 and -3 represent transcript regions 5 prime and 3 prime respectively)
DKFZP564O123 protein
opiate receptor-like 1
chloride channel, calcium activated, family member 2
v-jes-1 Y-enaguchi sarcoma viral oncogene homolog 1
coagulation factor IX (plasma thromboplastic component, Christmas disease, hemophilia B)
integrin, alpha 2 (CD49B, alpha 2 subunit of VLA-2 receptor)
nuclear factor IIC (CCAAT-binding transcription factor)
lipocalin-interacting membrane receptor
hypothetical protein FLJ10209
Consensus includes gb:AU152579 /FEA=EST /DB_XREF=gi:11014100 /DB_XREF=est:AU152579 /CLONE=NT2RP3001232 /UG=Hs.208931 Homo sapiens cDNA FLJ130 AU152579
Consensus includes gb:R73554 /FEA=EST /DB_XREF=gi:8417586 /DB_XREF=est:y9a05.x1 /CLONE=IMAGE:156416 /UG=Hs.103391 Human insulin-like growth factor bi R73554
phospholipase A2, group IIA (platelets, synovial fluid)
HCF-binding transcription factor Zhangfei
A1762013

DKFZP434F1735 protein
 KIAA1109 protein
 karyopherin alpha 2 (RAG cohort 1, importin alpha 1)
 Consensus includes gb:AU1417182 /FEA=EST /DB_XREF=est:AU1417182 /CLONE=HEMBB1002623 /UG=Hs.301197 Homo sapiens cDNA FLJ121471
 conserved gene amplified in osteosarcoma
 phosphoinositide-specific phospholipase C-beta 1
 hypothetical protein FLJ23129
 hypothetical protein FLJ20216
 Consensus includes gb:AL049259.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564E193 (from clone DKFZp564E193) /FEA=mRNA /DB_XREF=gi:4500005 /UG=Hs.11051
 adenylosuccinate lyase
 PRO1693 protein
 eukaryotic translation initiation factor 1A, Y chromosome
 ribonucleotide reductase M2 polypeptide
 catechol-O-methyltransferase
 phosphoribosyl pyrophosphate amidotransferase
 palmitoyl-protein thioesterase 2
 alcohol dehydrogenase 8 (class V)
 Consensus includes gb:AA837026 /FEA=EST /DB_XREF=gi:2912225 /DB_XREF=est:og09c12.s1 /CLONE=IMAGE:1439350 /UG=Hs.163549 ESTs
 low density lipoprotein receptor (familial hypercholesterolemia)
 excision repair cross-complementing rodent repair deficiency, complementation group 5 (xeroderma pigmentosum, complementation group G (Cockayne syndrome))
 erythropoietin
 Consensus includes gb:R94644 /FEA=EST /DB_XREF=gi:970039 /DB_XREF=est:yq42a12.1 /CLONE=IMAGE:198430 /UG=Hs.306542 Homo sapiens versican Vint isoform R94644
 prothymosin, alpha (gene sequence 28)
 hypothetical protein FLJ11042
 keratin, hair, acidic.2
 Ketch motif containing protein
 ring finger protein 25
 hypothetical protein FLJ11068
 phosphorylase kinase, alpha 1 (muscle)
 RAD51-like 1 (S. cerevisiae)
 HYA22 protein
 2,4-dienoyl CoA reductase 2, peroxisomal
 hypothetical protein FLJ10097
 hypothetical protein FLJ10560
 striated muscle contraction regulatory protein
 talin 2
 Consensus includes gb:AJ275355 /DEF=Homo sapiens partial IGVH3 V3-20 gene for immunoglobulin heavy chain V region, case 1, clone 2 /FEA=CDS /DB_XREF=gi:757; AJ275355
 glucocorticoid receptor DNA binding factor 1
 calpain 11
 glucagon
 phospholipase C, delta 1
 hypothetical protein FLJ23138
 integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)
 laminin, gamma 2 (nicotin (100KD), kalinin (105KD), B1M600 (100KD), Herlitz junctional epidermolysis bullosa))
 NCK adaptor protein 1
 LIM domain binding 1
 PAS-serine/threonine kinase
 IL-17B receptor
 cyclin G associated kinase
 A kinase (PRKA) anchor protein 10
 secretory carrier membrane protein 1
 AOP-ribosylation factor-like 3
 collagen, type I, alpha 1
 Down syndrome critical region gene 1-like 1
 KIAA0582 protein
 RAB5C, member RAS oncogene family
 KIAA0945 protein
 chitinase, di-N-acetyl-
 hypothetical protein FLJ21628
 mal transforming oncogene (derived from cell line NK14)- RAB8 homolog
 RAB6C, member RAS oncogene family
 phosphodiesterase 4D interacting protein (myomegalin)

melanoma antigen, family A, 4
cyclin-E binding protein 1
activin A receptor, type 1
winged-helix nude
geminin
hypothetical protein FLJ10595
Consensus includes gb:AL049593 /DEF=Human DNA sequence from clone RP4-654A7 on chromosome 20. Contains the gene for phospholipase C-beta-1 (KIAA0581, 1-p AL049593)
Consensus includes gb:AL137798 /DEF=Human DNA sequence from clone RPS-1182A14 on chromosome 1. Contains part of a gene similar to rat Espin, a pseudogene in AL137798
polydendylate binding protein-interacting protein 1
MAD, mothers against decapentaplegic homolog 9 (Drosophila)
KIAA0172 protein
BSP-like protein 2
v-kit Hardy-Zuckerman 4 feline sarcoma viral oncogene homolog
aldehyde dehydrogenase 3 family, member A2
growth factor receptor-bound protein 10
hypothetical protein DKFZP434K1210
transmembrane protein vezatin
zinc finger protein 143 (clone pHz-1)
DKFZP586F1018 protein
hypothetical protein FLJ11149
proprotein convertase subtilisin/kexin type 1
transducer of ERBB2, 2
phospholipase C, epsilon
KIAA0513 gene product
Consensus includes gb:AK000185.1 /DEF=Homo sapiens cDNA FLJ20178 fs, clone COL09990. /FEA=mRNA /DB_XREF=gi:7020104 /UG=Hs.306389 Homo sapiens cDNA AK000185
Huntingtin interacting protein H
DKFZP434M154 protein
deleted in lymphocytic leukemia, 2
Consensus includes gb:BC002465.1 /DEF=Homo sapiens, clone IMAGE:3344506. mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3344506) /DB_XI BC002465
Mad4 homolog
TU3A protein
Consensus includes gb:AF688610 /FEA=EST /DB_XREF=gi:4827918 /DB_XREF=est:y56a05.x5 /CLONE=IMAGE:162224 /UG=Hs.32043 ESTs
surfactant protein B
growth factor receptor-bound protein 10
KIAA0924 protein
dynein, cytoplasmic, intermediate polypeptide 1
Wolf-Hirschhorn syndrome candidate 1-like 1
Consensus includes gb:AF055018.1 /DEF=Homo sapiens clone 24442 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3005745 /UG=Hs.139137 Homo sapiens clone 24442 AF055018
PXR2b protein
Consensus includes gb:AL050204.1 /DEF=Homo sapiens mRNA; cDNA DKFZP586F1223 (from clone DKFZP586F1223). /FEA=mRNA /DB_XREF=gi:4884443 /UG=Hs.28: BF515597
KIAA1005 protein
lethal (3) malignant brain tumor (3jmbt) protein (Drosophila) homolog
protein (peptidyl-prolyl isomerase) NIMA-interacting, 4 (parvulin)
RAD23 homolog B (S. cerevisiae)
hypothetical protein FLJ20456
solute carrier family 13 (sodium/sulfate symporters), member 4
zinc finger protein 83 (HPF1)
BLUB1 budding uninhibited by benzimidazoles 1 homolog (yeast)
Consensus includes gb:AL109684.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 27080. /FEA=mRNA /DB_XREF=gi:5689805 /UG=Hs.306329 JAL109684
Consensus includes gb:X55989 /DEF=Human ECRP gene for eosinophil cationic related protein /FEA=mRNA /DB_XREF=gi:22940 /UG=Hs.166037 Human ECRP gene for X55989
Kruppel-like factor 1 (erythroid)
T-cell leukemia/lymphoma 6
weakly similar to glutathione peroxidase 2
insulin-like growth factor 1 (somatomedin C)
FK506 binding protein 8 (38kD)
flavin containing monooxygenase 3
hypothetical protein MGCA701
atriatin, calmodulin binding protein
pyruvate dehydrogenase kinase, isoenzyme 2
KIAA0599 protein
growth arrest-specific 7
hypothetical protein FLJ12975
Consensus includes gb:AL109681.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 112333. /FEA=mRNA /DB_XREF=gi:5689785 /UG=Hs.226017 AL109681

putative ribonuclease III	NM_013235
potassium voltage-gated channel, Shab-related subfamily, member 2	AF338730
DKFZP564O0823 protein	NM_015393
AXL receptor tyrosine kinase	AA679116
Consensus includes gb:A378044 /FEA=EST /DB_XREF=est:ie67h08.x1 /CLONE=IMAGE:2091809 /UG=Hs.23703 ESTs	A378044
microtubule-associated protein like ectoderm ENAP	AF103939
hypothetical protein DKFZP564A176	AL136663
WW domain binding protein 11	NM_016312
seizure related 6 homolog (mouse)-like	NM_021115
palmelphin	NM_017734
mitochondrial ribosomal protein S11	AL109953
hypothetical protein FLJ10058	NM_017885
ivb (bacterial acetylcholine synthase)-like	BC000109
DEME-6 protein	BC004399
homo box HB9	A173862
solute carrier family 25 (mitochondrial carrier: dicarboxylate transporter), member 10	NM_012140
Consensus includes gb:AF070628.1 /DEF=Homo sapiens clone 24803 /FEA=mRNA /DB_XREF=gi:3283895 /UG=Hs.32768 Homo sapiens clone 24803	AF070628
ivb (bacterial acetylcholine synthase)-like	A1680780
Nedd4-like ubiquitin-protein ligase	AL136902
hypothetical protein DKFZP434E169	BF447246
KIAA0060 protein	AA243659
hypothetical protein DKFZP566A1524	NM_025043
hypothetical protein FLJ22404	NM_029515
Consensus includes gb:AK024915.1 /DEF=Homo sapiens cDNA: FLJ1265 fls, done COL01484, /FEA=mRNA /DB_XREF=gi:10437330 /UG=Hs.306727 Homo sapiens cDNA: AK024915	NM_002108
histidine ammonia-lyase	AC002544
cardiolipin phosphatase, neuronal 3, juvenile (Batten, Spielmeier-Vogt disease)	AW299700
Consensus includes gb:AW299700 /FEA=EST /DB_XREF=est:es42p04.x1 /CLONE=IMAGE:2772342 /UG=Hs.18653 ESTs	NM_024231
Consensus includes gb:AK024231.1 /DEF=Homo sapiens cDNA FLJ14189 fls, done NT27RP2002058, /FEA=mRNA /DB_XREF=gi:10436558 /UG=Hs.306670 Homo sapiens cDNA: AK024231	AL023553
phosphomannomutase 1	NM_003140
sex determining region Y	NM_003140
programmed death ligand 2	NM_023238
protease, serine, 12 (neurolysin, motopsin)	NM_003619
Consensus includes gb:AV721430 /FEA=EST /DB_XREF=gi:10818582 /DB_XREF=est:AV721430 /CLONE=HTBBIF09 /UG=Hs.173638 transcription factor 7-like 2 (T-cell	AV721430
KIAA0373 gene product	NM_014684
cell death-inducing DFFA-like effector a	NM_001279
kidney and liver protein oxidase 1	U95090
proteasome beta 12	NM_018932
cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 6	NM_000782
transcription factor NRF	NM_017544
methylmalonate-semialdehyde dehydrogenase	AF130089
KIAA0234 protein	AW151108
proline-rich Glu (G-carboxyglutamic acid) polypeptide 2	NM_000951
phosphorylase, glycogen; muscle (McArdle syndrome, glycogen storage disease type V)	NM_005609
phosphodiesterase 9A	NM_002608
Wolff-Hirschhorn syndrome candidate 1-like 1	NM_023034
proline-rich protein BstNI subfamily 3	NM_006249
uncharacterized bone marrow protein BM040	NM_018456
Consensus includes gb:A086530 /FEA=EST /DB_XREF=gi:3424953 /DB_XREF=est:oz72a04.x1 /CLONE=IMAGE:1680846 /UG=Hs.255683 ESTs	A086530
major histocompatibility complex, class II, DR beta 4	U70544
keratin associated protein 4.7	AJ406939
mutated in colorectal cancers	NM_002387
myelin transcription factor 2	NM_003871
KIAA0758 protein	BF841489
Consensus includes gb:AK021569.1 /DEF=Homo sapiens cDNA FLJ11507 fls, done HEMBA1002160, /FEA=mRNA /DB_XREF=gi:10432775 /UG=Hs.314347 Homo sapiens cDNA: AK021569	AK021569
hemoglobin, beta	AF349114
Consensus includes gb:AW772192 /FEA=EST /DB_XREF=est:m69102.x1 /CLONE=IMAGE:3033147 /UG=Hs.7888 Homo sapiens clone 23736 m6	AW772192
mitogen-activated protein kinase kinase kinase 9	AF251442
tetraspan transmembrane 4 super family	NM_006675
hypothetical protein FLJ12541 similar to Sir6	AF352728
peptidyl arginine deiminase, type V	NM_012387
ribosomal protein L5	U66589
hypothetical protein FLJ11117	NM_018328
myelin protein zero (Charcot-Marie-Tooth neuropathy 1B)	U0537
secretogranin II (chromogranin C)	NM_003469

oxygen regulated protein (150KD)	NM_006389	1.26
transcription factor (p38 interacting protein)	A4889653	1.26
protein with polyglutamine repeat; calcium (ca2+) homeostasis endoplasmic reticulum protein	A372979	1.26
KIAA1919 protein	AK000168	1.26
melanoma antigen, family A, 8	BC002455	1.26
cytochrome b-5	M22978	1.26
CGI-75 protein	NM_016020	1.26
prostaglandin E receptor 2 (subtype EP2), 53kD	NM_000956	1.26
Consensus includes gb:AA195936 /FEA=EST /DB_XREF=gi:1791527 /DB_XREF=est:zp99a04.s1 /CLONE=IMAGE:628327 /UG=Hs.82719 Homo sapiens mRNA; cDNA DI AA195936	NM_000956	1.26
KIAA1324 protein	AB037745	1.26
ubiquitin specific protease 12	A820101	1.26
hypothetical protein FLJ10450	NM_018095	1.26
TAF10 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 30 kD	NM_006284	1.26
hypothetical protein FLJ12328	NM_024787	1.26
Consensus includes gb:AV700338 /FEA=EST /DB_XREF=gi:10302309 /DB_XREF=est:AV700338 /CLONE=GKCAMH12 /UG=Hs.286243 ESTs	AV700338	1.26
kelch-like ECH-associated protein 1	NM_012289	1.26
acetylserotonin O-methyltransferase	NM_004043	1.26
lipase, hepatic	NM_000236	1.26
homolog of Yeast RRP4 (ribosomal RNA processing 4), 3'-5'-exoribonuclease	BC000747	1.26
TNF receptor-associated factor 1	NM_005658	1.26
renal tumor antigen	NM_014226	1.26
hypothetical protein FLJ22318	NM_022762	1.26
potassium voltage-gated channel, KQT-like subfamily, member 1	AF051426	1.26
lymphocyte cytosolic protein 2 (SH2 domain-containing leukocyte protein of 76kD)	NM_005565	1.26
sialic acid binding Ig-like lectin 6	NM_001245	1.26
Consensus includes gb:AV468201 /FEA=EST /DB_XREF=gi:7038307 /DB_XREF=est:ha3a04.x1 /CLONE=IMAGE:2920878 /UG=Hs.91375 Human clone 23614 mRNA	NM_001245	1.26
nuclear phosphoprotein similar to S. cerevisiae PWP1	BE796924	1.26
RNA isopentenylpyrophosphate transferase	AF116679	1.26
camitine palmitoyltransferase L, liver	NM_001876	1.26
Consensus includes gb:AK024871.1 /DEF=Homo sapiens cDNA: FLJ21218 fs, clone COL00537, /FEA=mRNA /DB_XREF=gi:10437282 /UG=Hs.306712 Homo sapiens cDNA AK024871	NM_001876	1.26
trophin	NM_016157	1.26
ADP-ribosyltransferase 3	U47054	1.26
reticulocalbin 1, EF-hand calcium binding domain	NM_002901	1.26
EH-domain containing 3	NM_014600	1.26
Consensus includes gb:AIB84814 /FEA=EST /DB_XREF=gi:5569978 /DB_XREF=est:w85e05.x1 /CLONE=IMAGE:2431664 /UG=Hs.21964 ESTs	NM_014600	1.26
melanoma antigen, family A, 12	A884814	1.26
gb.L14457.1 /DEF=Human Ig rearranged kappa-chain gene V-J-region, complete cds, /FEA=mRNA /DB_XREF=gi:398491 /FL=gb.L14457.1	U82671	1.26
amine oxidase, copper containing 2 (retine-specific)	L14457	1.26
histamine N-methyltransferase	NM_009590	1.26
hypothetical protein PRO1777	NM_006895	1.26
KIAA0671 gene product	NM_018805	1.26
Consensus includes gb:AU145203 /FEA=EST /DB_XREF=gi:11006724 /DB_XREF=est:AU145203 /CLONE=HEMBA1004193 /UG=Hs.300897 Homo sapiens cDNA FLJ11111 AU145203	NM_018805	1.26
tenascin XB	BE044614	1.26
AU RNA binding protein/enoyl-Coenzyme A hydratase	NM_001698	1.26
secreted phosphoprotein 2, 24kD	NM_006944	1.26
fibroblast growth factor 23	NM_020638	1.26
zinc finger protein 313	AL031685	1.26
hypothetical protein MGC5528	L11372	1.26
annexin A7	NM_004034	1.26
megakaryocyte-enhanced gene transcript 1 protein	NM_021246	1.26
olfactory receptor, family 3Z, subfamily A, member 1	NM_012375	1.26
KIAA0196 gene product	NM_014846	1.26
phosphoglycerate kinase 1	AL121974	1.26
Consensus includes gb:AF090101.1 /DEF=Homo sapiens clone IMAGE 30714, /FEA=mRNA /DB_XREF=gi:4063636 /UG=Hs.177837 Homo sapiens clone IMAGE 30714	AL121974	1.26
G protein-coupled receptor 39	AF090101	1.26
kalikrein B, plasma (Fletcher factor) 1	AL567378	1.26
Consensus includes gb:AU141794 /FEA=EST /DB_XREF=gi:11008715 /DB_XREF=est:AU141794 /CLONE=HEMBA1002684 /UG=Hs.300978 Homo sapiens cDNA FLJ12	NM_000892	1.26
ADP-ribosylation factor 3	AW147184	1.26
calcium/calmodulin-dependent protein kinase (CaM kinase) II alpha	BQ341806	1.26
zinc finger protein 177	AW131813	1.26
Fancx1 anemia, complementation group G	NM_003451	1.26
tumor necrosis factor receptor superfamily, member 4	AC004472	1.26
olfactory receptor, family 12, subfamily D member 3	NM_003327	1.26
	NM_030959	1.26

zinc finger protein 286
hypothetical protein similar to preferentially expressed antigen of melanoma
erythropoietin receptor
SMC4 structural maintenance of chromosomes 4-like 1 (yeast)
membrane-spanning 4-domains, subfamily A, member 3 (hematopoietic cell-specific)
putative chemokine receptor; GTP-binding protein
KIAA0408 gene product
KIAA1750 protein
sperm acrosome associated 1
agrin
desmocollin 2
docking protein 1, 52KD (downstream of tyrosine kinase 1)
hypothetical protein MGC2827
matrix metalloproteinase 1 (interstitial collagenase)
dual specificity phosphatase 10
solute carrier family 2 (facilitated glucose transporter), member 10
hypothetical protein FLJ11017
KIAA0535 gene product
immunoglobulin mu binding protein 2
UDP glycosyltransferase 2 family, polypeptide B15
kinesin family member 1C
Consensus includes gb:BF939689 /FEA=EST /DB_XREF=gi:12357100 /DB_XREF=est:nac80b04.x1 /CLONE=IMAGE:3440575 /UG=Hs.159629 myosin IXB
DKF ZP564M1462 protein
KIAA1079 protein
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 12
chloride channel 2
Consensus includes gb:AK025072.1 /DEF=Homo sapiens cDNA: FLJ21419 fls, clone COL04084 /FEA=mRNA /DB_XREF=gi:10437512 /UG=Hs.306759 Homo sapiens cDNA: AK025072
radial spokehead-like 1
matrix metalloproteinase 23A
pellino homolog 2 (Drosophila)
brain abundant, membrane attached signal protein 1
laminin, gamma 1 (formerly LAMB2)
contactin 1
met proto-oncogene (hepatocyte growth factor receptor)
hypothetical protein FLJ10895
myelin proteolipid protein-like protein
phorbolin (similar to apolipoprotein B mRNA editing protein)
nact (homologous to the EG-AP (UBE3A) carboxyl terminus) domain and RCC1 (CHC1)-like domain (RLD) 1
deoxyribonuclease II, lysosomal
mitochondrial elongation factor G
translin
syndecan binding protein (synterin)
TEK tyrosine kinase, endothelial (venous malformations, multiple cutaneous and mucosal)
chromobox homolog 8 (Pc class homolog, Drosophila)
Chediak-Higashi syndrome 1
solute carrier family 12 (potassium/chloride transporters), member 4
guanine nucleotide binding protein (G protein), alpha inhibiting activity polypeptide 3
dual specificity phosphatase 3 (vaccinia virus phosphatase VH1-related)
collagen, type IX, alpha 1
I-complex 10 (a murine top homolog)
caldesmon 2 (cardiac muscle)
HERV-H LTR-associating 1
transducin-like enhancer of split 1 (Esf1) homolog, Drosophila
Consensus includes gb:AF070587.1 /DEF=Homo sapiens clone 24741 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3387960 /UG=Hs.25770 Homo sapiens clone 24741, AF070587
activity-regulated cytoskeleton-associated protein
periplakin
suppressor of variegation 3-9 (Drosophila) homolog 2; hypothetical protein FLJ23414
hBOIT for potent brain type organic ion transporter
uncoupling protein 1 (mitochondrial, proton carrier)
Consensus includes gb:AU150943 /FEA=EST /DB_XREF=gi:11012464 /DB_XREF=est:AU150943 /CLONE=NT2RP2003984 /UG=Hs.66762 Homo sapiens mRNA: cDNA (AU150943)
minor histocompatibility antigen HA-1

S-phase response (cyclin-related)
keratin, hair, acidic, 4
apoptosis signaling protein, nonapoptotic homolog (mouse)
Bacillus subtilis REF=M24537 /DEF=B subtilis pHeB, pheA genes corresponding to nucleotides 2897-3200 of M24537 /LEN=1409 (-5, -M, -3 represent transcript regions 5
M24537
potassium inwardly-rectifying channel, subfamily J, member 12
fetuin B
carboxylesterase 2 (intestine, liver)
arginase, liver
DC11 protein
KIAA0265 protein
Consensus includes gb:F09448 /FEA=EST /DB_XREF=gi:681981 /DB_XREF=est:HSC31G032 /CLONE=c-31g03 /UG=Hs.12840 Homo sapiens germline mRNA sequence
F09448
ata variant gene 3
apical protein-like (Xenopus laevis)
caspase 6, apoptosis-related cysteine protease
transmembrane 9 superfamily member 1
B-cell CLL/lymphoma 2
c-myc proto-oncogene tyrosine kinase
KIAA0557 protein
Consensus includes gb:AW452357 /FEA=EST /DB_XREF=gi:6993133 /DB_XREF=est:U1-H-B13-aii-4-03-Q-U1.s1 /CLONE=IMAGE:3068213 /UG=Hs.27373 Homo sapiens
M12874
estrogen receptor 1
Consensus includes gb:BE217875 /FEA=EST /DB_XREF=gi:8905193 /DB_XREF=est:hv31a03.x1 /CLONE=IMAGE:3174988 /UG=Hs.12311 Homo sapiens clone 23570 m
BE217875
sodium channel, voltage-gated, type II, beta polypeptide
pyruvate carboxylase
secreted ficolin-related protein 4
Consensus includes gb:AK000645.1 /DEF=Homo sapiens cDNA FLJ20628 fls, clone KAT02982, highly similar to SUI5_RABIT SUCRASE-ISOMALTASE /FEA=mRNA ID
AK000645
zinc finger protein 288
EGF-containing fibulin-like extracellular matrix protein 2
transglutaminase 3 (E polypeptide, protein-glutamine-gamma-glutamyltransferase)
exostosin (multiple) 1
intracellular hyaluronan-binding protein
hypothetical protein MGC3046
POU transcription factor
solute carrier family 8 (sodium-calcium exchanger), member 2
solute carrier family 22 (organic anion transporter), member 7
cytidine deaminase
microtubule-associated protein 7
Consensus includes gb:AB051440.1 /DEF=Homo sapiens mRNA for KIAA1653 protein, partial cds. /FEA=mRNA /GEN=KIAA1653 /PROD=KIAA1653 protein /DB_XREF=gi
AB051440
mitogen-activating protein kinase kinase kinase 2
progestin induced protein
gb:NM_017910.2 /DEF=Homo sapiens hypothetical protein FLJ20628 (FLJ20628), mRNA. /FEA=mRNA /GEN=FLJ20628 /PROD=hypothetical protein FLJ20628 /DB_XREF=gi
NM_017910
factor H-related protein 5
splicing factor, arginine/serine-rich 7 (35kD)
hypothetical protein FLJ22041 similar to FK506 binding proteins
cholinergic receptor, nicotinic, alpha polypeptide 9
Consensus includes gb:AA898858 /FEA=EST /DB_XREF=gi:3015728 /DB_XREF=est:ak23d05.s1 /CLONE=IMAGE:1406793 /UG=Hs.118567 ESTs
potassium voltage-gated channel, shaker-related subfamily, beta member 1
Rhesus blood group, CcEa antigens
small nuclear RNA activating complex, polypeptide 1, 43kD
mucin 16
cytidin-dependent kinase inhibitor 1C (p57, Kip2)
DKFZP434F122 protein
Consensus includes gb:AF339785.1 /DEF=Homo sapiens clone IMAGE:1963178, mRNA sequence. /FEA=mRNA /DB_XREF=gi:13507323 /UG=Hs.270256 Homo sapiens
cytochrome P450, subfamily 48 (cholesterol 24-hydroxylase)
hypothetical protein FLJ12810
hypothetical protein FLJ13511
protein tyrosine phosphatase, receptor type, N polypeptide 2
hormone box A5
soluble adenylyl cyclase
T-box and WD-40 domain protein 3
cytochrome b-561
Htn-1
Consensus includes gb:BF739971 /FEA=EST /DB_XREF=gi:12066647 /DB_XREF=est:7a41107.x1 /CLONE=IMAGE:3576732 /UG=Hs.98416 ESTs
BF739971

mitochondrial capsule selenoprotein 1.25 NM_030663

chromosome 11 open reading frame 11 1.25 AB014559

protein phosphatase 2 (formerly 2A), catalytic subunit, beta isoform 1.25 NM_004156

protocadherin gamma subfamily C, 3 1.25 AF152524

RIG-like 5-6 1.25 NM_015882

gb:BC006441.1 /DEF=Homo sapiens, Similar to RNA polymerase I transcription factor RRN3, clone MGC:13169, mRNA, complete cds. /FEA=mRNA /PROD=Similar to RN 1.25 BC006441

poly(IG) binding protein 2 1.25 AW103422

hypothetical protein DKFZp762E1312 1.25 NM_018410

distal-less homeobox 4 1.25 NM_001934

EphB2 1.25 AF025304

Consensus includes gb:AV741657 /FEA=EST /DB_XREF=gi:10859238 /DB_XREF=est:AV741657 /CLONE=CBMALG01 /UG=Hs.154085 leucine zipper protein 1 1.25 AV741657

chromodomain helicase DNA binding protein 4 1.25 NM_001273

prostacyclin D2 receptor (DP) 1.25 AI460323

cell division cycle 25C 1.25 AF277724

TAK1-binding protein 2 1.25 AL117407

hypothetical protein FLJ20366 1.25 NM_017786

mitogen-activated protein kinase-activated protein kinase 5 1.25 NM_003668

gb:AF116659.1 /DEF=Homo sapiens PRO1412 mRNA, complete cds. /FEA=mRNA /PROD=PRO1412 /DB_XREF=gi:7959816 /UG=Hs.321151 Homo sapiens PRO1412 m 1.25 AF116659

heterogeneous nuclear ribonucleoprotein L 1.25 X16135

KIAA0427 gene product 1.25 NM_014772

keratin 4 1.25 X07895

leucine-zipper-like transcriptional regulator, 1 1.25 NM_008767

insulin promoter factor 1, homeodomain transcription factor 1.25 U35632

solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7 1.25 NM_020309

NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2 (14.5kD, B14.5b) 1.25 NM_004549

p53-induced protein 1.25 NM_006034

hypothetical protein 1.25 NM_013341

Consensus includes gb:AW242315 /FEA=EST /DB_XREF=gi:6576069 /DB_XREF=est:xm96d12.x1 /CLONE=IMAGE:2692055 /UG=Hs.27860 Homo sapiens mRNA; cDNA 1.25 AW242315

peanut-like 2 (Orsophilis) 1.25 AF176379

interleukin 18 binding protein 1.25 NM_005699

hypothetical protein FLJ10312 1.25 NM_030672

hypothetical protein 1.25 NM_016629

KIAA0171 gene product 1.25 NM_014866

Wilms tumor associated protein 1.25 NM_015855

microfibrillar-associated protein 2 1.25 AL049569

hypothetical protein MGC3047 1.25 AW888223

Consensus includes gb:AL049690 /DEF=Human DNA sequence from clone 1080K8 on chromosome 20p12.1-13 Contains a pseudogene similar to 40S ribosomal protein 1.25 AL049690

hypothetical protein 1.25 W72694

amyloid beta (A4) precursor protein-binding, family A, member 3 (X11-like 2) 1.25 AI141541

Consensus includes gb:AL137495.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434C2331 (from clone DKFZp434C2331) /FEA=mRNA /DB_XREF=gi:5808119 /UG=Hs.30 1.25 AL137495

SRY (sex determining region Y)-box 11 1.25 AI360875

KIAA0579 protein 1.25 AL117532

chromosome 1 open reading frame 1 1.25 NM_001213

cholesterol receptor, muscarinic 2 1.25 NM_000739

ATPase, H⁺ transporting, lysosomal (vacuolar proton pump), alpha polypeptide, 70kD, isoform 1 1.25 AF113129

hypothetical protein FLJ10849 1.25 AI539361

protein tyrosine phosphatase, non-receptor type 22 (lymphoid) 1.25 NM_012411

reticulon 4 1.25 AF333338

fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome) 1.25 NM_023110

centaurin, beta 1 1.25 NM_014716

myeloproliferative leukemia virus oncogene 1.25 M90103

eukaryotic translation initiation factor 3, subunit 10 (theta, 150/170kD) 1.25 NM_003750

adenomedullin 1.25 NM_001124

olfactory receptor, family 7, subfamily E, member 12 pseudogene 1.25 AF065854

major histocompatibility complex, class II, DQ beta 1 1.25 AW276186

Fe fragment of IgG, low affinity IIa, receptor for (CD32) 1.25 NM_021642

pregnancy specific beta-1-glycoprotein 3 1.25 NM_021016

protein C (inactivator of coagulation factors Va and VIIIa) 1.25 NM_000312

solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2 1.25 AF064484

DEAD-box protein 1.25 NM_018665

adaptor-related protein complex 1, gamma 1 subunit 1.25 BF752277

KIAA0903 protein 1.25 BF116032

PDZ domain containing guanine nucleotide exchange factor(GEF)1 1.25 NM_014247

protein tyrosine phosphatase, non-receptor type 2 1.25
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease) 1.25
bradykinin receptor B1 1.25
G protein-coupled receptor 2 1.25
PBX/invited 1 homeobox 2 1.25
Cluster Ind. N53536;yz26n0.1.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:284209 /clone_end=3 /gb=N53536 /gi=1194702 /ug=Hs.33540 /len=543 1.25
ring finger protein 1 1.25
adenosine monophosphate deaminase (isoform E) 1.25
hypothetical protein FLJ22127 1.25
ubiquitin associated protein 1.25
lymphoid blast crisis oncogene 1.25
hypothetical protein FLJ22955 1.25
Mid-1-related chloride channel 1 1.25
glycerol kinase pseudogene 1 1.25
gb:BC000077.1 /DEF=Homo sapiens, Similar to KIAA0377 gene product, clone MGC:3288, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0377 gene product 1.25
a disintegrin and metalloproteinase domain 21 1.25
hypothetical protein FLJ23093 1.25
mut. homolog 3 (E. coli) 1.25
T-cell leukemia/lymphoma 6 1.25
tumor protein D52-like 1 1.25
KIAA0602 protein 1.25
likely ortholog of maternal embryonic leucine zipper kinase 1.25
tubby like protein 2 1.25
a disintegrin and metalloproteinase domain 29 1.25
fragile X mental retardation, autosomal homolog 1 1.25
kynureninase (L-kynurenine hydrolase) 1.25
phosphodiesterase 4A, cAMP-specific (phosphodiesterase E2 dunce homolog, Drosophila) 1.25
hypothetical protein DKFZ547M236 1.25
Consensus includes gb:AL050043.1 /DEF=Homo sapiens mRNA; cDNA DKFZ566M0524 (from clone DKFZ566M0524). /FEA=mRNA /DB_XREF=gi:4884285 /UG=Hs.3C /AL050043 1.25
gamma-glutamyltransferase 1 1.25
hypothetical protein FLJ23186 1.25
chromosome 21 open reading frame 108 1.25
cone-rod homeobox 1.25
integrin, beta 4 1.25
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1) 1.25
transcription factor AP-2 gamma (activating enhancer binding protein 2 gamma) 1.25
potassium inwardly-rectifying channel, subfamily J, member 1 1.25
plasminogen activator, urokinase 1.25
apolipoprotein B mRNA editing enzyme, catalytic polypeptide-like 2 1.25
leukocyte cell-derived chemotaxin 2 1.25
Consensus includes gb:AL037069 /FEA=EST /DB_XREF=gi:4073996 /DB_XREF=est:q82h04.x1 /CLONE=IMAGE:2009047 /UG=Hs.17518 Homo sapiens cDNA, par A1337069 1.25
cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 2 1.24
parathyroid hormone-like hormone 1.24
aniraxin A3 1.24
hypothetical protein FLJ10998 1.24
prostaglandin E receptor 4 (subtype EP4) 1.24
myosin VA (heavy polypeptide 12, myosin) 1.24
cerebellin 1 precursor 1.24
Consensus includes gb:BE223030 /FEA=EST /DB_XREF=gi:5910348 /DB_XREF=est:hu46c02.x1 /CLONE=IMAGE:3173090 /UG=Hs.301281 Homo sapiens mRNA, chroff BE223030 1.24
hydroxyacid oxidase 2 (long chain) 1.24
chromosome 8 open reading frame 1 1.24
VPS10 domain receptor protein 1.24
Consensus includes gb:AK024944.1 /DEF=Homo sapiens cDNA: FLJ21291 fts, clone COL01963. /FEA=mRNA /DB_XREF=gi:10437366 /UG=Hs.287657 Homo sapiens cDNA: AK024944 1.24
LIM homeobox protein 1 1.24
soluble carrier family 35 (UDP-galactose transporter), member 2 1.24
aristaeless-like homeobox 3 1.24
hypothetical protein PR02214 1.24
lymphocyte antigen 117 1.24
carbonic anhydrase XIV 1.24
protein kinase, cAMP-dependent, catalytic, alpha 1.24
dishevelled, dish homolog 2 (Drosophila) 1.24
GTPase regulator associated with the focal adhesion kinase pp125 1.24
Consensus includes gb:AB051438.1 /DEF=Homo sapiens mRNA for KIAA1651 protein, partial cds. /FEA=mRNA /GEN=KIAA1651 /PROD=KIAA1651 protein /DB_XREF=gi:AB051438 1.24

organic cation transporter OKB1	AL050350	1.24
hook 1 protein	NM_015888	1.24
hypothetical protein FLJ20555	NM_017877	1.24
guanine nucleotide binding protein (G protein) alpha 12	AK024696	1.24
Consensus includes gb:AL133576.1/DEE=Homo sapiens mRNA; cDNA DKFZp434L0850 (from clone DKFZp434L0850); partial cds. /FEA=mrRNA /GEN=DKFZp434L0850	AL133576	1.24
Consensus includes gb:BF447105/FEA=EST /DB_XREF=est/749607.x1 /CLONE=IMAGE:3649189 /UG=Hs.281706 sortlin 1	BF447105	1.24
Fc fragment of IgG, low affinity Iib, receptor for (CD32)	M31933	1.24
Consensus includes gb:AV733950 /FEA=EST /DB_XREF=gi:10851495 /DB_XREF=est/AV733950 /CLONE=cdAAADG12 /UG=Hs.326035 early growth response 1 /FL=gb:M	AV733950	1.24
TAF1 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 250 kD	M73444	1.24
ribosomal protein S6 kinase, 90kD, polypeptide 5	AF074393	1.24
v-akt murine thymoma viral oncogene homolog 3 (protein kinase B, gamma)	NM_005465	1.24
caplatin resistance associated	NM_006687	1.24
caldesmon 1 (fast-twitch, skeletal muscle)	NM_001231	1.24
ubiquitin-conjugating enzyme E2B (RAD8 homolog)	BC005979	1.24
glucosidase, alpha: acid (Pompe disease, glycogen storage disease type II)	NM_000152	1.24
KIAA1128 protein	AI478879	1.24
hypothetical protein PRO1163	NM_018576	1.24
hypothetical protein MGC5487	BC001305	1.24
myxovirus (influenza) resistance 2, homolog of murine	NM_002463	1.24
killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 1	NM_014512	1.24
ATP synthase, H ⁺ -transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 2	D13119	1.24
activin A receptor, type IB	NM_020327	1.24
inactive progesterone receptor, 23 kD	BE903880	1.24
phosphodiesterase 4C, cAMP-specific (phosphodiesterase E1 duncce homolog, Drosophila)	U83712	1.24
hypothetical protein FLJ14249 similar to HS1 binding protein 3	NM_022460	1.24
myotubularin related protein 7	AF073482	1.24
RAB35, member RAS oncogene family	NM_008861	1.24
nuclear receptor coactivator 2	AW027474	1.24
fibronogen, A alpha polypeptide	NM_000508	1.24
cell death-inducing DFFA-like effector b	NM_014430	1.24
nephirin related gene	NM_018240	1.24
unnamed HERV-H protein	U88896	1.24
phospholipase A2, group IVC (cytosolic, calcium-independent)	AF065214	1.24
beta-site APP-cleaving enzyme 2	NM_012105	1.24
endoplasmic reticulum resident protein 58; hypothetical protein MGC5302	NM_024089	1.24
cysteine and glycine-rich protein 3 (cardiac LIM protein)	NM_003476	1.24
t-complex 1	BF224073	1.24
cystic fibrosis transmembrane conductance regulator, ATP-binding cassette (sub-family C, member 7)	W60595	1.24
peroxisome proliferative activated receptor, gamma, coactivator 1	NM_013261	1.24
connexin46.6	BF061637	1.24
glutamate receptor, metabotropic 6	NM_000843	1.24
glycoprotein (transmembrane) nmb	NM_002510	1.24
glutathione S-transferase theta 1	NM_000853	1.24
small inducible cytokine subfamily B (Cys-X-Cys), member 14 (BRAK)	NM_004887	1.24
mitogen-activated protein kinase kinase kinase 12	BE222201	1.24
chromosome 8 open reading frame 1	BF057458	1.24
tripartite motif-containing 28	BE967532	1.24
UDP-Gal-beta-GlcNAc beta 1,3-galactosyltransferase, polypeptide 3	AB050656	1.24
ribosomal protein L5	AL137858	1.24
B-cell CLL/lymphoma 9	NM_004326	1.24
oculocerebrorenal syndrome of Lowe	NM_000276	1.24
ATPase, H ⁺ /K ⁺ exchanging, alpha polypeptide	NM_000704	1.24
gamma-glutamyltransferase 1	L20492	1.24
hepatocyte nuclear factor 3, beta	AI693985	1.24
erythrocyte nuclear factor 2, subfamily B, member 2	AJ302584	1.24
erythrocyte membrane protein band 7.2 (stomatrin)	AF130103	1.24
nuclear matrix protein p84	NM_005131	1.24
hypothetical protein from clone 643	BC003379	1.24
calcium/calmodulin-dependent protein kinase (CaM kinase) II beta	AF078803	1.24
BCL2-related protein A1	NM_004048	1.24
hypothetical protein FLJ13892	NM_025017	1.24
pervalbumin	NM_002854	1.24
sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6A	NM_020796	1.24

thyroid hormone receptor interactor 10	NM_004240	1.24
hypothetical protein MGC10327	TS4159	1.24
spinocerebellar ataxia 2 (olivopontocerebellar ataxia 2, autosomal dominant, ataxin 2)	NM_002973	1.24
leukocyte receptor cluster (LRC) member 4	NM_024298	1.24
deoxycyclidine kinase	NM_000788	1.24
Consensus includes gb:AK027254.1 /DEF=Homo sapiens cDNA: FLJ23601 f1s, clone LNG15501. /FEA=mRNA /DB_XREF=gi:10440343 /UG=Hs.306918 Homo sapiens cDNA: AK027254	AC002481	1.24
homologous to yeast nitrogen permease (candidate tumor suppressor)	NM_002177	1.24
interferon, omega 1	NM_004072	1.24
chemokine-like receptor 1	NM_013959	1.24
neuregulin 1	A053830	1.24
C-terminal binding protein 1	NM_006903	1.24
inorganic pyrophosphatase	NM_014622	1.24
loss of heterozygosity, 11, chromosomal region 2, gene A	AI753638	1.24
cardinal binding protein-like 2	AL049925	1.24
DKFZP547G0910 protein	BE973887	1.24
Consensus includes gb:AW975117 /FEA=EST /DB_XREF=gi:8166322 /DB_XREF=est:EST387223 /UG=Hs.145501 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAM1 AW975117	NM_018067	1.24
hairy homolog (Drosophila)	NM_018235	1.24
hypothetical protein FLJ10350	NM_024939	1.24
hypothetical protein FLJ10830	BC005153	1.24
hypothetical protein FLJ21918	NM_000780	1.24
reggulin 3A-like (without C2 domains)	X87870	1.24
cytochrome P450, subfamily VIIA (cholesterol 7 alpha-monooxygenase), polypeptide 1	NM_000512	1.24
hepatocyte nuclear factor 4, alpha	AJ300481	1.24
galactosamine (N-acetyl)-D-sulfate sulfatase (Morquio syndrome, mucopolysaccharidosis type IVA)	AF072164	1.24
chromosome 11 open reading frame 25	NM_005074	1.24
HFSE-1 protein	NM_001195	1.24
solute carrier family 17 (sodium phosphate), member 1	NM_001502	1.24
beaded filament structural protein 1, flensin	NM_005069	1.24
single-minded homolog 2 (Drosophila)	NM_001559	1.24
glycoprotein 2 (zymogen granule membrane)	NM_023067	1.24
interleukin 12 receptor, beta 2	NM_008464	1.24
forkhead box L2	BC006454	1.24
trans-golgi network protein 2	AK025116	1.24
growth arrest-specific 7	NM_002870	1.24
hypothetical protein FLJ21483	NM_016116	1.24
RAB13, member RAS oncogene family	NM_014448	1.24
ankyrin repeat and SOCS box-containing 4	NM_006561	1.24
Rho guanine exchange factor (GEF) 16	AF118663	1.24
CUG triplet repeat, RNA binding protein 2	AF110314	1.24
maternally expressed 3	NM_003489	1.24
Consensus includes gb:AF035307.1 /DEF=Homo sapiens clone 23785 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2861068 /UG=Hs.184697 Homo sapiens clone 23785 AF035307	AC002398	1.24
Consensus includes gb:AL117549.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586H0123 (from clone DKFZp586H0123). /FEA=mRNA /DB_XREF=gi:5912083 /UG=Hs.26 AL117549	BE138825	1.24
poliovirus receptor-related 1 (herpesvirus entry mediator C; nectin)	AI571208	1.24
regenerating islet-derived 1 alpha (pancreatic stone protein, pancreatic thread protein)	NM_014095	1.24
nuclear receptor interacting protein 1	M68013	1.24
hypothetical protein F25965	NM_002837	1.24
MCF-2 cell line derived transforming sequence-like	AF113683	1.24
major histocompatibility complex, class I, F	NM_030896	1.24
C18B11 homolog (44.9kD)	W28800	1.24
PRO1600 protein	BE856374	1.24
guanine nucleotide binding protein (G protein), alpha 11 (Gq class)	NM_024005	1.24
protein tyrosine phosphatase, receptor type, B	NM_017586	1.24
Consensus includes gb:AF113683.1 /DEF=Homo sapiens clone FLB3521. /FEA=mRNA /DB_XREF=gi:5855610 /UG=Hs.306569 Homo sapiens clone FLB3521	AL136480	1.24
hypothetical protein FLJ13520	NM_016079	1.24
Consensus includes gb:AF141901.1 /DEF=Homo sapiens GKAPSAPAP interacting protein (SHANK) mRNA, partial cds. /FEA=mRNA /GEN=SHANK /PROD=GKAPSAPAP AF141901	AW474615	1.24
Consensus includes gb:W28800 /FEA=EST /DB_XREF=gi:1308811 /DB_XREF=est:52g12 /UG=Hs.169802 ESTs	NM_013839	1.24
hypothetical protein FLJ12552		
DEAD(H) (Asp-Glu-Ala-Asp/His) box polypeptide 3		
chromosome 9 open reading frame 7		
proliferation-associated 204, 30kD		
CGI-149 protein		
Consensus includes gb:AW474615 /FEA=EST /DB_XREF=gi:7044721 /DB_XREF=est:xy18c02.x1 /CLONE=IMAGE:2653506 /UG=Hs.250988 ESTs		
olfactory receptor, family 10, subfamily H, member 2		

sialidase 3 (membrane sialidase)	NM_006656	1,24
glycerol kinase	AA292874	1,24
KIAA0060 protein	AA628780	1,24
potassium inwardly-rectifying channel, subfamily J, member 8	BF514158	1,24
MBIP protein	NM_016586	1,24
doublecortin and Cam kinase-like 1	NM_004734	1,24
leucine zipper-EF-hand containing transmembrane protein 1	NM_012318	1,24
histone deacetylase 8	NM_014707	1,24
Ca11	NM_019617	1,24
squamous cell carcinoma antigen recognized by T cell	NM_013352	1,24
interferon induced transmembrane protein 3 (1-8U)	BF338947	1,24
tubulin-specific chaperone d	AW247323	1,24
hypothetical protein FLJ13954	NM_025020	1,24
ryn-related kinase	NM_002031	1,24
putative GTP-binding protein similar to RAY/RAB1C	AL037167	1,24
KIAA0053 gene product	D29642	1,24
hypothetical protein FLJ10565	NM_018140	1,24
mitochondrial ribosomal protein S19C	NM_016067	1,24
KIAA1193 protein	AW003889	1,24
lipopolysaccharide binding protein	NM_004139	1,24
KIAA1048 protein	NM_014911	1,24
carbamoyl-phosphate synthetase 1, mitochondrial	W80357	1,24
patched related protein translocated in renal cancer	AF064801	1,24
glycine receptor, beta	NM_000824	1,24
protein tyrosine phosphatase, non-receptor type 3	NM_002829	1,24
erythroid differentiation-related factor	NM_016633	1,24
actin-like 6	NM_016188	1,24
Consensus includes gb:BE15346 /FEA=EST /DB_XREF=gi:9722561 /DB_XREF=est:601235986F1 /CLONE=IMAGE:3608205 /UG=Hs.278871 ESTs, Weakly similar to S-	BE15346	1,24
POB3B, member RAS oncogene family	BC005035	1,24
ribosomal protein S28	BF431363	1,24
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phosphoglycerate mutase 2 (muscle)	NM_000290	1,24
nitric oxide synthase 1 (neuronal)	NM_000620	1,24
intersectin 2	AF182199	1,24
angiopoietin-like 2	AI074333	1,24
hypothetical protein FLJ12789	NM_022495	1,24
homeo box (expressed in ES cells) 1	U82811	1,24
epithelial membrane protein 1	NM_001423	1,24
Consensus includes gb:AL049301.1 /DEF=Homo sapiens mRNA: cDNA DKFp564P073 (from clone DKFp564P073). /FEA=mRNA /DB_XREF=gi:4500087 /UG=Hs.2785	AL049301	1,24
mucin 1, transmembrane	AI810889	1,24
hypothetical protein FLJ23604	NM_025084	1,24
upstream binding protein 1 (LBP-1a)	NM_014517	1,24
MCF-2 cell line derived transforming sequence-like	AB002360	1,24
DNA segment, single copy probe LNS-CA/LNS-CAII (deleted in polyposis	BC000232	1,24
Wiskott-Aldrich syndrome-like	BE504979	1,24
short-chain alcohol dehydrogenase family member	NM_005794	1,24
chorea acanthocytosis	BF573874	1,24
KIAA0828 protein	AK025372	1,24
fibroblast growth factor 2 (basic)	M27968	1,24
Consensus includes gb:AA156777 /FEA=EST /DB_XREF=gi:1728392 /DB_XREF=est:18608.s1 /CLONE=IMAGE:502286 /UG=Hs.25431 KIAA1219 protein	AA156777	1,24
lysosomal serylase-like protein 1	NM_020354	1,24
chromosome condensation protein G	NM_022346	1,24
S100-type calcium binding protein A14	NM_020672	1,24
potassium inwardly-rectifying channel, subfamily J, member 16	NM_018658	1,24
telomerase reverse transcriptase	NM_003219	1,24
DEADH (Asp-Glu-Ala-AspHis) box polypeptide 28	AI822937	1,24
hypothetical protein FLJ10743	NM_018201	1,24
interleukin 26	NM_018402	1,24
connector enhancer of KSR-like (Drosophila kinase suppressor of ras)	NM_008314	1,24
B1 for mucin	X83412	1,24
hypothetical protein	BF572868	1,24
RNA guanylyltransferase and 5'-phosphatase	NM_003800	1,24
aspartyl aminopeptidase	AI690563	1,24

	AK025938	hypothetical protein
	NM_001166	baculoviral IAP repeat-containing 2
	AF230389	tpramide motif-containing 2B
	AV657604	krefelin A-inhibited guanine nucleotide-exchange protein 2
	NM_003820	fucose-1-phosphate guanylyltransferase
	AF064084	isopenicillamine carboxyl methyltransferase
	BE312885	leucine zipper, putative tumor suppressor 1
	NM_001150	alaninyl (membrane) aminopeptidase N, aminopeptidase M, microsomal aminopeptidase, CD(13, p150)
	NM_004163	RAB27B, member RAS oncogene family
	ZN_AAT69008	Consensus includes gb:AAT69008 /FEA=EST /DB_XREF=gj:582080c10.s1 /CLONE=IMAGE:1318578 /UG=Hs.290921 EST's, Highly similar to G0ZT AA769008
	NM_000341	solute carrier family 3 (cycline, dibasic and neutral amino acid transporters, activator of cycline, basic and neutral amino acid transport), member 1
	NM_Z072475	Consensus includes gb:A072475 /FEA=EST /DB_XREF=gj:5769391 /DB_XREF=est:w4d0b09.x1 /CLONE=IMAGE:2490137 /UG=Hs.243501 Homo sapiens cDNA FLJ227475
	AK025938	FSGS17
	BF217301	transducin-like enhancer of split 4 (E(spl)) homolog, Drosophila
	NM_006103	WAP four-disulfide core domain 2
	AI478592	DNA-dependent protein kinase catalytic subunit-interacting protein 2
	AF153419	Inhibitor of kappa light polypeptide gene enhancer in B-cells, kinase complex-associated protein
	AL047886	hypothetical protein CGO18
	U24078	killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 1
	AB020682	F-box only protein 21
	AL133620	likely homolg of rat kinase D-interacting substance of 220 kDa
	U82457	glutamate receptor, metabotropic 4
	NM_017879	collagen, type VI, alpha 3
	NM_004369	solute carrier family 21 (organic anion transporter), member 9
	AF205073	Consensus includes gb/AK021571.1 /DEF=Homo sapiens cDNA FLJ11509 flt, clone HEMBA1002166, /FEA=mRNA /DB_XREF=gj:10432777 /UG=Hs.289093 Homo sapie AK021571
	NM_001871	analogouscoprotein rector 1
	AB059408	hypothetical protein SMP31
	BC002710	kallikrein 10
	AF230629	dihydropyrimidinase-like 3
	NM_001387	C2 torf19-like protein
	NM_015955	HRA5-like supressor
	NM_020386	hypothetical protein FLJ11082
	NM_018317	keratin, cuticle, ultrahigh sulphur 1
	NM_005553	leucine zipper-EF-hand containing transmembrane protein 1
	AJ359368	US zinner-associated Sm-like protein
	AJ153477	CGI-147 protein
	NM_018077	coactivator-associated arginine methyltransferase-1
	AA551784	T cell receptor beta locus
	AF009787	deleted in azoospermia 4
	AF248483	Rev/Rex activation domain binding protein-related
	NM_006076	I/K cytokine, down-regulator of HLA II
	AF182645	sevelin
	AF796813	Consensus includes gb/AK024606.1 /DEF=Homo sapiens cDNA: FLJ20953 fls, clone ADSEDI079, /FEA=mRNA /DB_XREF=gj:10436920 /UG=Hs.306897 Homo sapiens cAK024606
	NM_004842	A thase (PRKA) anchor protein 7
	NM_004944	deoxyribonucleas e-Like 3
	AU148222	hephaestin
	NM_002060	gap junction protein, alpha 4, 37kD (connxin 37)
	AA534894	Cluster intn, AA534894/i82d11.a1 Homo sapiens cDNA, 3 end /clone=IMAGE-926421 /clone_end=3 /gb=AA534894 /gj=2279147 /ug=Hs.4914 /len=602
	AV721013	hypothetical protein IMAGE:4215339
	NM_002060	Consensus includes gb:AY242701 /FEA=EST /DB_XREF=gj:8576546 /DB_XREF=est:m9a0a0x.1 /CLONE=IMAGE:26891446 /UG=Hs.256398 Homo sapiens mRNA; cdN AY242701
	NM_004891	rmyeloidysplasia syndrome 1
	AI935096	nuclear receptor subfamily 4, group A, member 2
	NM_004794	RAB33A, member RAS oncogene family
	BFG69886	Consensus includes gb:BFG69886 /FEA=EST /DB_XREF=gj:12337201 /DB_XREF=gj:12337201 /DB_XREF=est:g0z272821f1 /CLONE=IMAGE:4360804 /UG=Hs.170226 Homo sapiens clone Z35: BFG69886 X61078
	X61078	T cell receptor alpha variable 8-B
	D28121	Consensus includes gb:D28121.1 /DEF-Human mRNA for ZFM1 protein alternatively spliced product, complete cds, /FEA=mRNA /PROD=ZFM1 protein, alternately spliced product /DB_XRE D28121
	AFDS9180	picyclin, basic protein of sperm head cystoskeleton 1
	Z22780	heterogeneous nuclear ribonucleoprotein D-like
	OBB9878	A thase (PRKA) anchor protein 2
	NM_003007	semenogelin 1
	NM_003007	

CGI-04 protein	NM_015936	1,23
hormo box A2	NM_006735	1,23
potassium channel, subfamily K, member 5 (TASK-2)	NM_003740	1,23
hypothetical protein FLJ11125	NM_018331	1,23
SMY3 suppressor of mit two 3 homolog 2 (yeast)	AI871724	1,23
hypothetical protein FLJ13231	NM_023073	1,23
winged-type MMTV integration site family, member 11	NM_004626	1,23
gap junction protein, alpha 7, 45KD (connexin 45)	NM_005497	1,23
KIAA0090 protein	BF187122	1,23
Consensus includes gb:AL066728.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D044 (from clone DKFZp434D044). /FEA=mRNA /DB_XREF=gi:5419863 /UG=Hs.1465	AL066728	1,23
Consensus includes gb:AK023895.1 /DEF=Homo sapiens cDNA FLJ13833 fls. clone THYR01000676 /FEA=mRNA /DB_XREF=gi:10435969 /UG=Hs.296745 Homo sapiens	AK023895	1,23
Consensus includes gb:AA401963 /FEA=EST /DB_XREF=gi:2055965 /DB_XREF=est:u53208.s1 /CLONE=IMAGE:741710 /UG=Hs.194107 ESTs. Moderately similar to AI	AA401963	1,23
mesothelin	NM_005823	1,23
Consensus includes gb:AI927479 /FEA=EST /DB_XREF=gi:5663443 /DB_XREF=est:wb90006.x1 /CLONE=IMAGE:2462603 /UG=Hs.165590 ribosomal protein S13	AI927479	1,23
EBNA1 binding protein 2	NM_006824	1,23
mitogen-activated protein kinase 9	W37431	1,23
T-box 5	NM_000182	1,23
Consensus includes gb:BE783279 /FEA=EST /DB_XREF=gi:10204477 /DB_XREF=est:601470835F.1 /CLONE=IMAGE:3873801 /UG=Hs.33862 ESTs	BE783279	1,23
BTB and CNC homology 1, basic leucine zipper transcription factor 1	AF026189	1,23
arradillo repeat gene deletion in velocardiofacial syndrome	NM_001670	1,23
semitr/SUMO-specific protease 3	AX000923	1,23
Consensus includes gb:AL043266 /FEA=EST /DB_XREF=gi:5935844 /DB_XREF=est:DKFZp434L1423.s1 /CLONE=DKFZp434L1423 /UG=Hs.111334 famitin, light polypep	AL043266	1,23
Consensus includes gb:BC002778.1 /DEF=Homo sapiens. Similar to myosin light chain 2, precursor lymphocyte-specific, clone MGC:3479, mRNA, complete cds. /FEA=mRNA /PROD=Similar	BC002778	1,23
solute carrier family 8 (neurotransmitter transporter, GABA), member 11	NM_014229	1,23
transcription factor binding to IGHM enhancer 3	AL181985	1,23
methionine sulfoxide reductase A	NM_012331	1,23
MRS2-like, magnesium homeostasis factor (S. cerevisiae)	AF052167	1,23
H1 histone family, member T (testis-specific)	NM_005323	1,23
Consensus includes gb:AI537540 /FEA=EST /DB_XREF=gi:4451675 /DB_XREF=est:tp04g11.x1 /CLONE=IMAGE:2188852 /UG=Hs.71168 Homo sapiens clone 24674 mR	AI537540	1,23
Consensus includes gb:AI979278 /FEA=EST /DB_XREF=gi:5804295 /DB_XREF=est:wb03304.x1 /CLONE=IMAGE:2515854 /UG=Hs.177462 ESTs. Weakly similar to KIAA	AI979278	1,23
KIAA0321 protein	AB002319	1,23
somatostatin	NM_001048	1,23
glycoprotein V (platelet)	L11238	1,23
CTAGE-1 protein	NM_022663	1,23
vasoactive intestinal peptide receptor 1	NM_004624	1,23
putative homeodomain transcription factor 1	NM_006608	1,23
KIAA0285 gene product	R61539	1,23
G protein-coupled receptor 12	NM_005286	1,23
putative N-acetyltransferase Camello 2	NM_016347	1,23
Consensus includes gb:AA045174 /FEA=EST /DB_XREF=gi:1523376 /DB_XREF=est:z66a07.s1 /CLONE=IMAGE:487764 /UG=Hs.16163 Homo sapiens mRNA; cDNA DI	AA045174	1,23
fatty-acyl-Coenzyme A ligase, long-chain 6	AV727634	1,23
Rho guanine exchange factor (GEF) 15	NM_014958	1,23
transglutaminase 4 (prostate)	NM_003241	1,23
Consensus includes gb:AL049337.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564P016 (from clone DKFZp564P016). /FEA=mRNA /DB_XREF=gi:4500118 /UG=Hs.1325	AL049337	1,23
N-acetylglucosamine 6-phosphatase (acid ceramidase)	AI379338	1,23
translocase of outer mitochondrial membrane 34	NM_006809	1,23
kallikrein 8 (neuropathicovasin)	NM_007196	1,23
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) membrane sector associated protein M8-9	NM_014471	1,23
serine protease inhibitor, Kazal type 4	AF248966	1,23
XO4803 B subunit thrc, thrb genes corresponding to nucleotides 248-2229 of XO4603 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	XO4603	1,23
trefoil factor 3 (intestinal)	NM_003226	1,23
phospholipase A2, group IID	NM_012400	1,23
carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 7	AL022165	1,23
fructosamine-3-kinase	NM_022158	1,23
Consensus includes gb:AL138548.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761G18121 (from clone DKFZp761G18121); complete cds. /FEA=mRNA /GEN=DKFZp761	AL138548	1,23
serum/glucoconoid regulated kinase 2	NM_016276	1,23
dynamin intermediate chain 2	AF250288	1,23
Consensus includes gb:AL080068.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564J062 (from clone DKFZp564J062). /FEA=mRNA /DB_XREF=gi:5262475 /UG=Hs.27253	AL080068	1,23
topoisomerase (DNA) III alpha	AI876092	1,23
Consensus includes gb:W87634 /FEA=EST /DB_XREF=gi:1401696 /DB_XREF=est:zh87d09.s1 /CLONE=IMAGE:417113 /UG=Hs.17947 ESTs. Weakly similar to K02F3.11	W87634	1,23
Jagged 2	AF029778	1,23
hypothetical protein MGC2376	AF872101	1,23
CGI-79 protein	AF078865	1,23

secreted phosphoprotein 1 (osteopontin, bone sialoprotein 1, early T-lymphocyte activation 1) 1,23
 tyrosinase-related protein 1 1,23
 lymphocyte antigen 117 1,23
 myosin IA 1,23
 lymphocyte antigen 117 1,23
 pleckstrin B2 1,23
 Cluster Ind. T87245:y91b02.r1 Homo sapiens cDNA, 5' and (clone=IMAGE:115583 /clone_end=5' /gb=T87245 /gi=715597 /ug=Hs.16336 /len=412 1,23
 leukocyte immunoglobulin-like receptor, subfamily A (with TM domain), member 1 1,23
 solute carrier family 7 (carbonic amino acid transporter, y+ system), member 4 1,23
 CD84 antigen (leukocyte antigen) 1,23
 hypothetical protein FLJ20837 1,23
 Consensus includes gb:AK024949.1 /DEF=Homo sapiens cDNA: FLJ21296 fls, clone COL02029 /FEA=mRNA /DB_XREF=gi:10437372 /UG=Hs.306734 Homo sapiens cDNA: AK024949 1,23
 calcium binding protein 1 (calbriain) 1,23
 organic cation transporter 1,23
 tumor necrosis factor receptor superfamily, member 21 1,23
 prostate differentiation factor 1,23
 kinesin family member 3A 1,23
 TAF9-like RNA polymerase II, TATA box binding protein (TBP)-associated factor, 31 kD 1,23
 Consensus includes gb:A081543 /FEA=EST /DB_XREF=gi:3416335 /DB_XREF=est:DKFZ56602446_s1 /CLONE=IMAGE:1555750 /UG=Hs.300851 ESTs, Highly similar to T0870 1,23
 B-factor, properdin 1,23
 parvin, beta 1,23
 midline 2 1,23
 translocase of inner mitochondrial membrane 10 homolog (yeast) 1,23
 hypothetical protein FLJ12180 1,23
 Consensus includes gb:AK025038.1 /DEF=Homo sapiens cDNA: FLJ21385 fls, clone COL03356 /FEA=mRNA /DB_XREF=gi:10437474 /UG=Hs.306750 Homo sapiens cDNA: AK025038 1,23
 Consensus includes gb:AL038824 /FEA=EST /DB_XREF=gi:5407955 /DB_XREF=est:DKFZ56602446_s1 /CLONE=DKFZ56602446 /UG=Hs.12840 Homo sapiens germ 1,23
 forthead box O3A 1,23
 solute carrier family 9 (sodium/hydrogen exchanger), isoform 6 1,23
 granzyme M (lymphocyte met-ase 1) 1,23
 neurotrophic tyrosine kinase, receptor, type 3 1,23
 death effector filament-forming Ced-4-like apoptosis protein 1,23
 likely ortholog of mouse p16i like homolog 1 (Orsophila)-like 1,23
 c-fos induced growth factor (vascular endothelial growth factor D) 1,23
 gonadotropin-releasing hormone receptor 1,23
 ubiquitin specific protease 19 1,23
 Consensus includes gb:A1478795 /FEA=EST /DB_XREF=gi:4373608 /DB_XREF=est:U52607.x1 /CLONE=IMAGE:2161741 /UG=Hs.41123 ESTs, Moderately similar to T1 1,23
 Consensus includes gb:A1346431 /FEA=EST /DB_XREF=gi:4083637 /DB_XREF=est:qp50g09.x1 /CLONE=IMAGE:1926466 /UG=Hs.75188 wee1+ (S. pombe) homolog 1,23
 phosphatidylinositol 4-kinase type II 1,23
 hypothetical protein FLJ20618 1,23
 phospholipid scramblase 1 1,23
 Consensus includes gb:AW974666 /FEA=EST /DB_XREF=gi:8165852 /DB_XREF=est:EST386755 /UG=Hs.293024 ESTs 1,23
 v-abl Abelson murine leukemia viral oncogene homolog 2 (arg, Abelson-related gene) 1,23
 mitogen inducible 2 1,23
 MAX protein 1,23
 hypothetical protein from EUROMAGE 511235 1,23
 EGF-containing fibulin-like extracellular matrix protein 1 1,23
 T-box 1 1,23
 KIAA0590 gene product 1,23
 Consensus includes gb:AW971248 /FEA=EST /DB_XREF=gi:8161093 /DB_XREF=est:EST383337 /UG=Hs.291289 ESTs, Weakly similar to ALU1_HUMAN ALU SUBFAM1 1,23
 chondroitin sulfate proteoglycan 5 (neuroglycan C) 1,23
 titin-cap (telethonin) 1,23
 Mas4 homolog 1,23
 adaptor-related protein complex 3, delta 1 subunit 1,23
 fatty acid binding protein 5 (psoriasis-associated) 1,23
 KIAA1046 protein 1,23
 hypothetical protein FLJ10534 1,23
 KIAA0140 gene product 1,23
 collagen, type V, alpha 3 1,23
 nuclear receptor subfamily 0, group B, member 1 1,23
 hypothetical protein MGC4655 1,23
 atrophin-1 interacting protein 1; activin receptor interacting protein 1 1,23
 transcription elongation factor B (SIII), polypeptide 2 (18kD, elongin B) 1,23
 hypothetical protein MGC5590 1,23

Rhesus blood group-associated glycoprotein 1.23
Down syndrome critical region gene 1 1.23
gb:BC000651.1 /DEF=Homo sapiens, Similar to solute carrier family 1 (glutamate transporter), member 7, clone MGC:2078, mRNA, complete cds. /FEA=mRNA /PROD=S- 1.23
HX44 antigen X91103 1.23
Consensus includes gb:AA576497 /FEA=EST /DB_XREF=gi:2353597 /DB_XREF=est:nm:7607.s1 /CLONE=IMAGE:1074133 /UG=Hs.252497 ESTs 1.23
hypothetical protein FLJ22639 1.23
nuclear factor of activated T-cells 5, tonically-responsive 1.23
protein kinase, cAMP-dependent, regulatory, type I, beta 1.23
KIAA0740 gene product 1.23
xylosylprotein beta1 (4-galactosyltransferase, polypeptide 7 (galactosyltransferase I) 1.23
KIAA0887 protein 1.23
Consensus includes gb:D60132 /FEA=EST /DB_XREF=gi:961771 /DB_XREF=est:HUM087G05A /CLONE=GEN-087G05 /UG=Hs.79581 Human clone 23560 mRNA seque 1.23
stromal cell-derived factor 2 1.23
non-metastatic cells 5, protein expressed in (nucleoside-diphosphate kinase) 1.23
BCL2-associated athanogene 3 1.23
integral membrane protein 2A 1.23
cortactin SH3 domain-binding protein 1.23
sialic acid binding Ig-like lectin 7 1.23
solute carrier family 4, anion exchanger, member 3 1.23
ATPase, Ca++ transporting, plasma membrane 3 1.23
tumor necrosis factor receptor superfamily, member 10d, decoy with truncated death domain 1.23
monokine induced by gamma interferon 1.23
very long-chain acyl-CoA synthetase homolog 2 1.23
Charot-Leyden crystal protein 1.23
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila) 1.23
flamogen-like 1 1.23
protein tyrosine phosphatase, receptor type, E 1.23
PBX/Ino101 1 homeobox 2 1.23
pleiotrophin (heparin binding growth factor 8, neurite growth-promoting factor 1) 1.23
ATPase, Ca++ transporting, cardiac muscle, fast twitch 1 1.23
Consensus includes gb:BE46371 /FEA=EST /DB_XREF=gi:5511146 /DB_XREF=est:hw15c12.x1 /CLONE=IMAGE:3182998 /UG=Hs.227170 ESTs 1.23
par-6 partitioning defective 8 homolog beta (C. elegans) 1.23
inter-alpha (globulin) inhibitor H4 (plasma kallikrein-sensitive glycoprotein) 1.23
putative selenocysteine lyase 1.23
iroquois homeobox protein 4 1.23
heterogeneous nuclear ribonucleoprotein A1 1.23
retinoic acid induced 3 1.23
chromosome 5 open reading frame 4 1.23
monoamine oxidase B 1.23
pregnancy-associated plasma protein A 1.23
protein kinase C, mu 1.23
neuromedin U 1.23
KIAA0009 gene product 1.23
meb-21-like 1 (C. elegans) 1.23
RNA, U17D small nucleolar 1.23
CD24 antigen (small cell lung carcinoma cluster 4 antigen) 1.23
olfactory receptor, family 4, subfamily F, member 3 1.23
lamtinh, gamma 3 1.23
schwannomin interacting protein 1 1.23
KIAA0885 protein 1.23
Consensus includes gb:AK023515.1 /DEF=Homo sapiens cDNA FLJ13453 fts, clone PLACE1003205. /FEA=mRNA /DB_XREF=gi:10435471 /UG=Hs.301715 Homo sapier 1.23
Kruppel-like factor 12 1.23
hypothetical protein PRO1483 1.23
uroplakin 1B 1.23
hypothetical protein PRO2176 1.23
phosphatidylinositol glycan, class A (paroxysmal nocturnal hemoglobinuria) 1.23
guanine nucleotide binding protein (G protein), alpha 14 1.23
sex hormone-binding globulin 1.23
Consensus includes gb:AL137547.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434B1120 (from clone DKFZp434B1120). /FEA=mRNA /DB_XREF=gi:6808230 /UG=Hs.25 AL137547 1.23
ATP-binding cassette, sub-family C (CFTR/MRP), member 8 1.23
asialoglycoprotein receptor 2 1.23

phosphomannomutase 2	NM_000303	1,23
Rai guanine nucleotide exchange factor RaiGPS1A	NM_014636	1,23
cytoplasmic FMRP interacting protein 1	BC005097	1,23
Consensus includes gb:A1351043 /FEA=EST /DB_XREF=est:q22a08.x1 /CLONE=IMAGE:1948310 /UG=Hs.279477 ESTs	A1351043	1,23
KIAA0913 protein	W69120	1,23
hypothetical protein FLB3342	NM_020118	1,23
scute carrier family 17 (sodium phosphate), member 2	NM_005835	1,23
DNA segment on chromosome 12 (unique) 2493 expressed sequence	NM_007360	1,23
calreticulin	AD000692	1,23
adenosine A2b receptor	NM_000676	1,23
mammaglobin 1	NM_002411	1,23
SP110 nuclear body protein	NM_004510	1,23
myeloproliferative leukemia virus oncogene	NM_005373	1,23
cadherin, EGF LAG seven-pass G-type receptor 2 (flamingo homolog, Drosophila)	D87469	1,23
hypothetical protein FLJ11539	NM_024748	1,23
fibroblast growth factor 18	NM_003862	1,23
immunoglobulin heavy constant mu	X95660	1,23
Consensus includes gb:AA654586 /FEA=EST /DB_XREF=gi:2590740 /DB_XREF=est:n0411.s1 /CLONE=IMAGE:1192173 /UG=Hs.325505 ESTs, Moderately similar to H	AA654586	1,23
a disintegrin and metalloproteinase domain 8	AB14527	1,22
caveolin 2	BF197655	1,22
teratocarcinoma-derived growth factor 1	NM_003212	1,22
cystatin C (amyloid angiopathy and cerebral hemorrhage)	NM_000099	1,22
src family associated phosphoprotein 1	NM_003726	1,22
gb:AF122828.1 /DEF=Homo sapiens neurofibromatosis type 2 protein isoform Mer151 (NF2) mRNA, alternatively spliced, complete cds. /FEA=mRNA (GEN=NF2 /PROD=	AF122828	1,22
Consensus includes gb:AV088547 /FEA=EST /DB_XREF=gi:6044358 /DB_XREF=est:xd17c08.x1 /CLONE=IMAGE:2594030 /UG=Hs.312076 ESTs, Moderately similar to	AV088547	1,22
BUB1 budding inhibited by benzimidazoles 1 homolog (yeast)	AK023540	1,22
peroxisome receptor 1	NM_000319	1,22
potassium voltage-gated channel, subfamily G, member 1	A132876	1,22
X-prolyl aminopeptidase (aminopeptidase P) 2, membrane-bound	NM_003399	1,22
immunoglobulin superfamily, member 4	NM_014333	1,22
histamine receptor H3	NM_007232	1,22
Consensus includes gb:AA773554 /FEA=EST /DB_XREF=gi:2825125 /DB_XREF=est:ab61a05.s1 /CLONE=IMAGE:845264 /UG=Hs.234216 Human 40871 mRNA partial s	AA773554	1,22
egfrn	AW008051	1,22
thymosin, beta, Identified in neuroblastoma cells	NM_021992	1,22
CD7 antigen (p41)	NM_006137	1,22
rng finger protein (C3HC4 type) 8	NM_003958	1,22
vascular Rho-GAP/TBC-containing	BC001663	1,22
CD36 antigen (collagen type I receptor, thrombospondin receptor)-like 2 (lysosomal integral membrane protein II)	NM_005506	1,22
hypothetical protein FLJ20534	NM_017867	1,22
Consensus includes gb:AL356504 /DEF=Human DNA sequence from clone RP1-14N1 on chromosome 1q21.1-21.3 Contains ESTs, GSSs and STSs. Contains the FLG ge	AL356504	1,22
zinc finger protein 181	BG035116	1,22
protease, serine, 8 (prolactin)	NM_002773	1,22
histidine-rich glycoprotein	NM_006412	1,22
KIAA0864 protein	AB020671	1,22
FOXJ2 forkhead factor	NM_018416	1,22
hemochromatosis	BG402460	1,22
calenin (cadherin-associated protein), delta 2 (neural plakophilin-related arm-repeat protein)	U96136	1,22
amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease)	BC004369	1,22
KIAA1644 protein	BF516072	1,22
hypothetical protein FLJ12969	NM_022638	1,22
Bacteriophage /REF=X03453 /DEF=Bacteriophage P1 cre recombinase protein corresponding to nucleotides 1032-1270 of X03453 /LEN=1058 /5 and -3 represent transcr	X03453	1,22
KIAA0800 gene product	NM_014703	1,22
tissue inhibitor of metalloproteinase 3 (Sorby fundus dystrophy, pseudoinflammatory)	U87195	1,22
Consensus includes gb:AF254357.1 /DEF=Homo sapiens prostate-specific membrane antigen PSM mRNA, exon 8 alternative splice variant, partial cds. /FEA=mRNA /PRI	AF254357	1,22
neurexophilin 4	AF933199	1,22
Consensus includes gb:AV712064 /FEA=EST /DB_XREF=gi:10731370 /DB_XREF=est:AV712064 /CLONE=DCAAUD05 /UG=Hs.9456 SWISNF related, matrix associated,	AV712064	1,22
myomesin 1 (skelemin) (185kD)	NM_003803	1,22
gb:NM_005196.1 /DEF=Homo sapiens centromere protein F (350400KD, miosin) (CENPF), mRNA. /FEA=mRNA (GEN=CENPF /PROD=centromere protein F (350400KD,	NM_005196	1,22
CGI-142	AK001280	1,22
Consensus includes gb:AL031118 /DEF=Human DNA sequence from clone 153G14 on chromosome 6p21.3-22.2. Contains the 3 end of the ZNF164 gene for Kruppel-like	AL031118	1,22
collagen, type IV, alpha 4	D17391	1,22
hypothetical protein FLJ12484	NM_022767	1,22
Consensus includes gb:AF009287.1 /DEF=Homo sapiens clone FBA1 C14-du-chat region mRNA. /FEA=mRNA /DB_XREF=gi:2331069 /UG=Hs.102238 Homo sapiens clone	AF009287	1,22

vascular cell adhesion molecule 1	NM_001078	1.22
displein resistance-associated overexpressed protein	AW089673	1.22
pp28, US snRNP 100 kd protein	AF028402	1.22
Consensus includes gb:AL133089.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434E2028 (from clone DKFZp434E2028) /FEA=mRNA /DB_XREF=gi:64533541 /UG=Hs.27	AL133089	1.22
CUG triplet repeat, RNA binding protein 2	N36839	1.22
run-related transcription factor 2	NM_004348	1.22
ADP-ribosylation factor 4	NM_001660	1.22
likely ortholog of mouse pwl like homolog 1 (Drosophila)-like	NM_018068	1.22
SCAN domain-containing 1	AL109965	1.22
major histocompatibility complex, class II, DP alpha 2 (pseudogene)	X01634	1.22
testis-specific kinase 2	NM_007170	1.22
immunoglobulin heavy constant mu	AB035175	1.22
kinase suppressor of ras	AA971768	1.22
DKFZP568N0721 protein	NM_015400	1.22
hypothetical protein FLJ14251	NM_024881	1.22
topoisomerase (DNA) II binding protein	NM_007027	1.22
DKFZP564F0522 protein	AL049943	1.22
hypothetical protein R29124_1	BF110434	1.22
Consensus includes gb:AL353943.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761E0611 (from clone DKFZp761E0611) /FEA=mRNA /GEN=DKFZp761E0611 /PROD=hy	AL353943	1.22
neurochondrin	AB011179	1.22
UL18 binding protein 1	NM_025218	1.22
cytoplasmic linker 2	BC006259	1.22
matrix Gla protein	AL532375	1.22
hypothetical gene CG018	AL049786	1.22
Consensus includes gb:AL307586 /FEA=EST /DB_XREF=gi:4002311 /DB_XREF=est:35h12.x1 /CLONE=IMAGE:2056391 /UG=Hs.133130 Homo sapiens mRNA; cDNA	AL307586	1.22
KIAA0417 gene product	AB007877	1.22
Consensus includes gb:AL535683 /FEA=EST /DB_XREF=gi:4449818 /DB_XREF=est:cong1.P11.A9 /UG=Hs.327729 ESTs	AL535683	1.22
EF4-like factor 5 (eta domain transcription factor)	NM_001422	1.22
CD1C antigen, c polypeptide	NM_001785	1.22
bromodomain adjacent to zinc finger domain, 2A	AI653128	1.22
v-erb-b2 erythroblastic leukemia viral oncogene homolog 3 (avian)	NM_001982	1.22
Consensus includes gb:AL109666.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMAGE 35907 /FEA=mRNA /DB_XREF=gi:5683800 /UG=Hs.7242 Ho	AL109666	1.22
hypothetical protein MGC12780	BC006312	1.22
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13	AJ001696	1.22
gbU19346.1 /DEF=Human (tpi-met fusion) oncogene mRNA, complete cds. /FEA=mRNA /GEN=tpi-met fusion /PROD=tpi-met fusion protein /DB_XREF=gi:625085 /FL=gb	U19346	1.22
Consensus includes gb:BF062139 /FEA=EST /DB_XREF=gi:10821049 /DB_XREF=est:7N72110.x1 /CLONE=IMAGE:3481059 /UG=Hs.282387 polymerase (RNA) III (DNA	BF062139	1.22
hypothetical protein DKFZp564D0372	NM_031301	1.22
olfactory receptor, family 2, subfamily H, member 3	NM_007160	1.22
Consensus includes gb:AK022038.1 /DEF=Homo sapiens cDNA FLJ11978 fls, clone HEMBB1001253 /FEA=mRNA /DB_XREF=gi:10433356 /UG=Hs.296679 Homo sapie	AK022038	1.22
hypothetical protein MGC4170	NM_024312	1.22
Notch homolog 2 (Drosophila)	NM_024408	1.22
nuclear receptor subfamily 1, group 1, member 3	NM_005122	1.22
claudin 18	NM_016369	1.22
hypothetical protein	AI005473	1.22
GW128 protein	NM_014052	1.22
T cell receptor beta locus	U66059	1.22
EphA5	X95425	1.22
SRY (sex determining region Y)-box 13	AF116571	1.22
Consensus includes gb:BG109555 /FEA=EST /DB_XREF=gi:12603061 /DB_XREF=est:60228068F1 /CLONE=IMAGE:4367968 /UG=Hs.331207 ESTs	BG109555	1.22
apoptotic protease activating factor	AF248734	1.22
GRB2-related adaptor protein 2	NM_004810	1.22
scellin	NM_003843	1.22
synovial sarcoma, X breakpoint 3	BF164274	1.22
KIAA0514 gene product	NM_014686	1.22
transmembrane protease, serine 2	NM_006656	1.22
Consensus includes gb:AF090886.4 /DEF=Homo sapiens clone HQ0072 /FEA=mRNA /DB_XREF=gi:6690153 /UG=Hs.284192 Homo sapiens clone HQ0072	AF090886	1.22
biotinidase	AI787414	1.22
Consensus includes gb:D16471.1 /DEF=Human mRNA, Xq terminal portion. /FEA=mRNA /DB_XREF=gi:495099 /UG=Hs.121571 Human mRNA, Xq terminal portion	D16471	1.22
hypothetical protein FLJ22844	NM_025098	1.22
taste receptor, type 2, member 4	NM_016944	1.22
M10098 Human 18S rRNA gene, complete (L5_M_3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	M10098	1.22
solute carrier family 4, anion exchanger, member 2 (erythrocyte membrane protein band 3-like 1)	NM_003040	1.22
gamma-glutamyltransferase-like activity 1	NM_004121	1.22

myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10
 signal-regulatory protein beta 2
 synovial sarcoma, X breakpoint 3
 small inducible cytokine subfamily A (Cys-Cys), member 23
 phospholipase A2-activating protein
 suppressor of Ty 3 homolog (S. cerevisiae)
 small proline-rich protein 2B
 collagen, type I, alpha 2
 tumor protein p53-binding protein
 solute carrier family 4, sodium bicarbonate cotransporter, member 7
 PRO1992 protein
 olfactory receptor, family 1, subfamily A, member 2
 hypothetical protein FLJ23590
 gamma-aminobutyric acid (GABA) A receptor, beta 1
 desmoplakin (DPI, DPII)
 hypothetical protein MGC4694
 carbonic anhydrase XII
 chromosome 5 open reading frame 4
 tubulin, beta polypeptide 4, member Q
 KIAA0853 protein
 Consensus includes gb:AV1715767 /FEA=EST /DB_XREF=gi:10797284 /DB_XREF=est:AV1715767 /CLONE=DCBATH02 /UG=Hs.7780 Homo sapiens mRNA; cDNA DKFZ: AV1715767
 transmembrane 4 superfamily member 1
 hemoglobin, alpha 2
 G protein-coupled receptor C5L2
 nuclear receptor coactivator 1
 potassium voltage-gated channel, KQT-like subfamily, member 4
 similar to rat HREY107
 C3H-type zinc finger protein; similar to D. melanogaster muscleblind B protein
 potassium voltage-gated channel, KQT-like subfamily, member 2
 fucosyltransferase 6 (alpha (1.3) fucosyltransferase)
 Consensus includes gb:AI188346 /FEA=EST /DB_XREF=gi:3739555 /DB_XREF=est:q08a10.x1 /CLONE=IMAGE:1723098 /UG=Hs.301778 ESTs, Weakly similar to ALU1 AI188346
 aladin 2-binding protein 1
 Consensus includes gb:AV720803 /FEA=EST /DB_XREF=gi:10817955 /DB_XREF=est:AV720803 /CLONE=GLCCMA04 /UG=Hs.293055 ESTs
 DNase II-like acid DNase
 carboxyl ester lipase-like (bile salt-stimulated lipase-like)
 brain-specific membrane-anchored protein
 KIAA0974 protein
 cyclin F
 mannan-binding lectin serine protease 2
 FGFR1 oncogene partner
 Consensus includes gb:BE881590 /FEA=EST /DB_XREF=gi:10330366 /DB_XREF=est:601490008F1 /CLONE=IMAGE:3892465 /UG=Hs.10684 Homo sapiens clone 2442
 sialyltransferase
 reticulation 4
 Consensus includes gb:AA370433 /FEA=EST /DB_XREF=gi:2344433 /DB_XREF=est:tnk6211.1.s1 /CLONE=IMAGE:1018125 /UG=Hs.10323 Homo sapiens mRNA from chr
 signal transduction protein (SH3 containing)
 calmodulin-like 3
 KIAA0603 protein
 Consensus includes gb:BG111761 /FEA=EST /DB_XREF=gi:12605267 /DB_XREF=est:602285343F1 /CLONE=IMAGE:4372619 /UG=Hs.8107 Homo sapiens mRNA; cDN
 serum/glucocorticoid regulated kinase 2
 hypothetical protein FLJ12929
 KIAA0492 protein
 Consensus includes gb:AK000257 1 /DEF=Homo sapiens cDNA FLJ20250 fs, clone COLF6635, /FEA=mRNA /DB_XREF=gi:7020217 /UG=Hs.274505 Homo sapiens mR
 PRO0618 protein
 EGF-like-domain, multiple 6
 H326
 Cyclin M2
 potassium inwardly-rectifying channel, subfamily J, member 10
 hypothetical protein
 type I intermediate filament cytoskeleton
 fibroblast growth factor 16
 guanylate binding protein 1, interferon-inducible, 67kD
 LIM homeobox protein 2
 cyclic nucleotide gated channel beta 1

NM_004841 1.22
 NM_018556 1.22
 BC005904 1.22
 U58913 1.22
 AF083395 1.22
 AF084804 1.22
 NM_008945 1.22
 NM_000089 1.22
 AF150117 1.22
 NM_003615 1.22
 NM_014107 1.22
 NM_012352 1.22
 NM_024649 1.22
 NM_000812 1.22
 NM_004415 1.22
 BC004911 1.22
 NM_001218 1.22
 H93077 1.22
 U83110 1.22
 BE895685 1.22
 AV715767 1.22
 AF189753 1.22
 V00489 1.22
 NM_018485 1.22
 BF576458 1.22
 NM_004700 1.22
 BC001387 1.22
 A088145 1.22
 D82346 1.22
 U27331 1.22
 NM_018723 1.22
 AV720803 1.22
 NM_021233 1.22
 NM_001808 1.22
 NM_012109 1.22
 AW241779 1.22
 NM_001761 1.22
 NM_006610 1.22
 NM_007045 1.22
 BE881590 1.22
 NM_006456 1.22
 AF320999 1.22
 AA570453 1.22
 NM_005864 1.22
 N38707 1.22
 N34407 1.22
 BG111761 1.22
 Z98752 1.22
 NM_024937 1.22
 R02172 1.22
 AK000257 1.22
 NM_014133 1.22
 NM_015507 1.22
 AU146233 1.22
 AF216962 1.22
 NM_002241 1.22
 NM_016617 1.22
 NM_015515 1.22
 NM_003868 1.22
 NM_002053 1.22
 NM_004789 1.22
 U58637 1.22

KIAA0657 protein
acyl-Coenzyme A dehydrogenase, long chain desmulin
Consensus includes gb:AK027128.1 /DEF=Homo sapiens cDNA: FLJ23475 /fs, clone HSI13659 /FEA=mRNA /DB_XREF=gi:10440177 /UG=Hs.288982 Homo sapiens cDI AK027128
DKFZP564G2022 protein
likely homolog of mouse immunity-associated nucleotide 4
transglutaminase 2 (C polypeptide, protein-glutamine-gamma-glutamyltransferase)
gb:BC006387.1 /DEF=Homo sapiens, clone MGC:12878, mRNA, complete cds. /FEA=mRNA /PROD=unknown (protein for MGC:12878) /DB_XREF=gi:13623518 /FL=gb:BC006387
M98478
AK002166
BE551361
NM_017545
BC002351
AL042842
D13665
NM_004829
NM_030765
NM_014297
Z76308
AF091086
AF281859
NM_014799
L78668
NM_006520
AK000089
NM_003157
NM_005253
AV705485
NM_017976
A854858
cDNA DI AA114843
Consensus includes gb:AA114843 /FEA=EST /DB_XREF=gi:1669911 /DB_XREF=est:20212.s1 /CLONE=IMAGE:491183 /UG=Hs.183454 Homo sapiens mRNA: cDNA DI AA114843
AF212842
NM_016582
NM_001486
NM_005481
NM_018659
BE674860
U88358
BG280646
NM_000576
NM_005641
NM_006068
U82164
NM_014965
U20428
NM_012368
NM_002758
AA479278
NM_004438
NM_004850
NM_009587
M18767
BC005236
AF055270
NM_000896
A823980
NM_012785
NM_012082
H79306
X55448
AK000851
N34846

split hand/foot malformation (ectrodactyly) type 3
hnpheasin
Consensus includes gb:AV706096 /FEA=EST /DB_XREF=gi:10723384 /DB_XREF=est:AV706096 /CLONE=ADBAWH11 /UG=Hs.326248 Homo sapiens cDNA: FLJ22071
killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 2
t-complex-associated-testis-expressed 1-like
HDCMA18P protein
serine/threonine kinase 2
FOS-like antigen 2
KIAA1038 protein
hypothetical protein FLJ10038
Plasivie prostate cancer tumor suppressor
Consensus includes gb:AA114843 /FEA=EST /DB_XREF=gi:1669911 /DB_XREF=est:20212.s1 /CLONE=IMAGE:491183 /UG=Hs.183454 Homo sapiens mRNA: cDNA DI AA114843
Consensus includes gb:AA114843 /FEA=EST /DB_XREF=gi:1669911 /DB_XREF=est:20212.s1 /CLONE=IMAGE:491183 /UG=Hs.183454 Homo sapiens mRNA: cDNA DI AA114843
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 7
peptide transporter 3
glucokinase (hexokinase 4) regulatory protein
v-maf musculoaponeurotic fibrosarcoma oncogene homolog B (avian)
cytokine-like protein C17
mutS homolog 5 (E. coli)
ubiquitin-conjugating enzyme E2N (UBC13 homolog, yeast)
Consensus includes gb:U88358.1 /DEF=Homo sapiens clone R2 EtbB-3 R2 (c-erbB-3) mRNA, partial cds. /FEA=mRNA /GEN=c-erbB-3 /PROD=ErB-3 R2 /DB_XREF=gi:108358
TAF5 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 80 kD
tol-like receptor 6
antigen identified by monoclonal antibodies 12E7, F21 and O13
KIAA1042 protein
suppression of tumorigenicity 14 (colon carcinoma, matrilysin, epithin)
olfactory receptor, family 2, subfamily C, member 1
mitogen-activated protein kinase kinase 6
Consensus includes gb:AA479278 /FEA=EST /DB_XREF=gi:2207834 /DB_XREF=est:z12107.s1 /CLONE=IMAGE:754261 /UG=Hs.9295 elastin (supravalvular aortic stenosis)
EphA4
dermatan sulfate proteoglycan 3
lectin, galactoside-binding, soluble, 9 (galectin 9)
complement component 1, s subcomponent
hypothetical protein MGC12262
heat-shock suppressed protein 1
cytochrome P450, subfamily IVF, polypeptide 3 (leukotriene B4 omega hydroxylase)
ribokinase
Consensus includes gb:AA127885 /FEA=EST /DB_XREF=gi:1687192 /DB_XREF=est:z13504.s1 /CLONE=IMAGE:501775 /UG=Hs.13350 Homo sapiens mRNA: cDNA DI AA127885
Friend of GATA2
platelet-derived growth factor receptor, alpha polypeptide
glucose-6-phosphate dehydrogenase
chromosome 22 open reading frame 4
DAZ associated protein 2

tocopherol (alpha) transfer protein (ataxia (Friedreich-like) with vitamin E deficiency)
 E2F transcription factor 3
 myosin, heavy polypeptide 3, skeletal muscle, embryonic
 lysozyme (oculocutaneous albinism A)
 protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (caldesmon A gamma)
 Consensus includes gb:AF032176.1 /DEF=Homo sapiens clone 24457 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360487 /UG=Hs.158529 Homo sapiens clone 24457
 glutamine-fructose-6-phosphate transaminase 2
 early lymphoid activation protein
 advanced glycosylation end product-specific receptor
 hypothetical protein FLJ13920
 cyclin-dependent kinase inhibitor 1C (p57, Kip2)
 pleckstrin homology-like domain, family A, member 1
 KIAA0934 protein
 transcription factor BMAL2
 kallikrein 2, prostatic
 beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase I)
 zinc finger protein 282
 chitinase 3-like 1 (cartilage glycoprotein-39)
 CGI-40 protein
 HMG-box transcription factor TCF-3
 Consensus includes gb:AK023774.1 /DEF=Homo sapiens cDNA FLJ13712 fls. clone PLACE2000394. /FEA=mRNA /DB_XREF=gi:10435808 /UG=Hs.273099 Homo sapiens
 carcinoembryonic antigen-related cell adhesion molecule 4
 collagen, type XVII, alpha 1
 KIAA0173 gene product
 chromosome 21 open reading frame 2
 hypothetical protein FLJ12729
 integrin, alpha 9
 solute carrier family 28 (sodium-coupled nucleoside transporter), member 1
 RAB1, member RAS oncogene family
 Consensus includes gb:AK001613.1 /DEF=Homo sapiens cDNA FLJ10757 fls. clone NT2RP3004578, highly similar to Homo sapiens mRNA for KIAA0477 protein. /FEA=m
 carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen)
 solute carrier family 18 (monocarboxylic acid transporters), member 2 (putative transporter)
 nuclear receptor co-repressor 1
 hypothetical protein MGCA293
 hypothetical protein FLJ10956
 sialyltransferase 8C (alpha2,3Galbeta1,4GlcNAcalpha2,8-sialyltransferase)
 programmed cell death 1
 olfactory receptor, family 11, subfamily A, member 1
 hypothetical protein FLJ11871
 sorting nexin 2
 Cluster Ind. A1801973:28605.x1 Homo sapiens cDNA, 3' end (clone=IMAGE:2270985 (clone_end=3' (gb=A1801973 (gi=5367445 /ug=Hs.27373 /len=532
 carboxylesterase 1 (monocysteine/macrophage serine esterase 1)
 Sjogren syndrome antigen A1 (SZKD, ribonucleoprotein autoantigen SS-A/Ro)
 methionine adenosyltransferase II, alpha
 activated p21cdc42Hs kinase
 prostaglandin E receptor 3 (subtype EP3)
 olfactory receptor, family 2, subfamily W, member 1
 SAR1 protein
 solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 3
 lactalbumin, alpha-
 Lutheran blood group (Auberger b antigen included)
 Lutheren blood critical region gene 3
 oculocutaneous syndrome of Lowe
 fatty acid binding protein 3, pseudogene 2
 serologically defined breast cancer antigen 84
 Consensus includes gb:AC004010 /DEF=Human BAC clone GS1-99H8 /FEA=COS /DB_XREF=gi:2781385 /UG=Hs.121520 Human BAC clone GS1-99H8
 solute carrier family 9 (sodium/hydrogen exchanger), isoform 5
 hypothetical protein PRD-NY3
 sarcoglycan, alpha (50KD dystrophin-associated glycoprotein)
 glutamate receptor, metabotropic 1
 RAB27A, member RAS oncogene family
 mucin 1, transmembrane

MHC binding factor, beta
CGI-35 protein
Zinc finger protein 2 heterolazy 1 (odd-paired homolog, Drosophila)
melanoma antigen, family A, 11
thyroid stimulating hormone receptor
eyes absent homolog 1 (Drosophila)
Tax interaction protein 1
hypothetical protein FLJ13857
hepocidin antimicrobial peptide
KIAA0547 gene product
Consensus includes gb:A198543 /FEA=EST /DB_XREF=gi:3751149 /DB_XREF=est:q49f11.x1 /CLONE=IMAGE:1753389 /UG=Hs.8982 ESTs, Highly similar to KIAA1395 A198543
sarcoglycan, delta (35KD dystrophin-associated glycoprotein)
POZ-75 protein
cytosolic beta-glucosidase
hypothetical protein FLJ22690
H4 histone family, member C
KIAA1219 protein
Consensus includes gb:BG536504 /FEA=EST /DB_XREF=gi:13528050 /DB_XREF=est:602564909F1 /CLONE=IMAGE:4689314 /UG=Hs.274518 Homo sapiens mRNA; cf BG536504
AL570914
PBX/Notch1 homeobox 1
mamaglobin 2
FAT tumor suppressor homolog 2 (Drosophila)
chondroitin sulfate proteoglycan BEHAB/brevican
cyclin-dependent kinase inhibitor 1B (p27, Kip1)
Consensus includes gb:BE985029 /FEA=EST /DB_XREF=gi:11768976 /DB_XREF=est:601658812R1 /CLONE=IMAGE:3886131 /UG=Hs.198793 Homo sapiens cDNA: FL BE985029
karyopherin beta 2b, transportin
insulin-like growth factor binding protein 6
soluble carrier family 6 (neurotransmitter transporter, creatine), member 8
hypothetical protein AF053356_CDS3
Werner syndrome
Consensus includes gb:AW979018 /FEA=EST /DB_XREF=gi:8170301 /DB_XREF=est:EST391128 /UG=Hs.132670 ESTs
l-mis domain-containing protein
dual specificity phosphatase 4
retinal homeobox protein
hypothetical protein EST00098
tumor necrosis factor (ligand) superfamily, member 11
KIAA0570 gene product
ATP binding protein associated with cell differentiation
chromosome 5 open reading frame 7
secreted protein, acidic, cysteine-rich (osteonectin)
protein tyrosine phosphatase, non-receptor type 1
KIAA1068 protein
wingless-type MMTV integration site family, member 5A
Consensus includes gb:AW404894 /FEA=EST /DB_XREF=gi:6923951 /DB_XREF=est:U1-HF-BL0-act-f07-0-U1.r1 /CLONE=IMAGE:3059844 /UG=Hs.314762 Homo sapiens mRNA; cf AW404894
glucosylase 1
major histocompatibility complex, class II, DQ beta 1
hypothetical protein MGC3035
lipin 2
Consensus includes gb:AW008921 /FEA=EST /DB_XREF=gi:5857699 /DB_XREF=est:wa75a03.x1 /CLONE=IMAGE:2503756 /UG=Hs.13138 Homo sapiens, clone IMAGE AW008921
KIAA0931 protein
Consensus includes gb:AF103295.1 /DEF=Homo sapiens clone N97 immunoglobulin heavy chain variable region mRNA, partial cds. /FEA=mRNA /PROD=immunoglobulin AF103295
KIAA1067 protein
hypothetical protein FLJ10853
CDC42 binding protein kinase beta (DMPK-like)
KIAA0938 protein
LIM and senescent cell antigen-like domains 1
APR-1 protein
guanylate cyclase 1, soluble, alpha 3
annexin A3
cofactor required for Sp1 transcriptional activation, subunit 6 (77KD)
TAF11 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 28 KD
N-methylpurine-DNA glycosylase
chromosome X open reading frame 2
ret proto-oncogene (multiple endocrine neoplasia and medullary thyroid carcinoma 1, Hirschsprung disease)

putative mitochondrial solute carrier
 prominin-like 1 (mouse)
 SRY (sex determining region Y)-box 13
 myosin, heavy polypeptide 7, cardiac muscle, beta
 hHOC for homolog of Drosophila headcase
 ankyrin 1, erythrocytic
 ankyrin 1, erythrocytic
 embryonic ectoderm development
 acetylserotonin O-methyltransferase-like
 hypothetical protein FLJ20706
 SH3 domain binding glutamic acid-rich protein like
 nebulin
 granzyme B (granzyme 2, cytotoxic T-lymphocyte-associated serine esterase 1)
 oligomannin 1
 islet cell autoantigen 1 (69KD)
 hypothetical protein
 T cell receptor gamma locus
 prostaglandin D2 synthase (21KD, brain)
 hypothetical protein FLJ13287
 hypothetical protein FLJ10084
 scd homology three (SH3) and cysteine rich domain
 UDP-glycosyltransferase 1 family, polypeptide A3
 proline synthetase co-transcribed (bacterial homolog)
 villin 1
 RNA (guanine-7-)-methyltransferase
 Consensus includes gb:AL512688.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547J2313 (from clone DKFZp547J2313) /FEA=mRNA /GEN=DKFZp547J2313 /PROD=hyp
 zinc finger protein 37 homolog (mouse)
 gb:NM_016154.2 /DEF=Homo sapiens hyaluronic acid receptor (HAR), mRNA /FEA=mRNA /GEN=HAR /PROD=hyaluronic acid receptor /DB_XREF=gi:10835259 /UG=Hs.NM_016164
 Fanconi anemia, complementation group G
 Consensus includes gb:AA558468 /FEA=EST /DB_XREF=gi:2328945 /DB_XREF=est:nc39c06.s1 /CLONE=IMAGE:1015882 /UG=Hs.178665 Homo sapiens immunoglobulin A558468
 MB02 (methyl-CpG-binding protein)-interacting zinc finger protein
 interferon, beta 1, fibroblast
 pyruvate dehydrogenase kinase, isoenzyme 2
 zinc finger protein 278
 tissue inhibitor of metalloproteinase 3 (Sorsby fundus dystrophy, pseudoinflammatory)
 hepatocyte growth factor (hepatopoietin A; scatter factor)
 chloride channel 6
 solute carrier family 24 (sodium/potassium/calcium exchanger), member 1
 Consensus includes gb:AK001442.1 /DEF=Homo sapiens cDNA FLJ10560 fs, clone NT2RP2003533 /FEA=mRNA /DB_XREF=gi:7022701 /UG=Hs.140402 Homo sapiens cDNA AK001442
 Consensus includes gb:AK028273.1 /DEF=Homo sapiens cDNA: FLJ22620 fs, clone HSI05629 /FEA=mRNA /DB_XREF=gi:10439077 /UG=Hs.306834 Homo sapiens cDNA AK028273
 killer cell lectin-like receptor subfamily B, member 1
 hypothetical protein FLJ12886
 solute carrier family 21 (organic anion transporter), member 14
 matrix metalloproteinase 13 (collagenase 3)
 fms-related tyrosine kinase 3 ligand
 forkhead box E2
 melastatin 1
 gonadotropin-releasing hormone receptor
 interleukin 10
 KIAA0599 protein
 frizzled homolog 9 (Drosophila)
 solute carrier family 16 (monocarboxylic acid transporters), member 1
 wingless-type MMTV integration site family, member 4
 heparanase-like protein
 insulin-like 3 (Leydig cell)
 hypothetical protein FLJ12983
 interleukin 1 receptor antagonist
 hypothetical protein DKFZp762P2111
 BCE-1 protein
 heart and neural crest derivatives expressed 1
 coagulation factor XII (Hageman factor)
 KIAA0752 protein
 SET binding factor 1

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potassium inwardly-rectifying channel, subfamily J, member 5
 HSPC063 protein
 serine protease inhibitor-like, with Kunitz and WAP domains 1 (epplin)
 family with sequence similarity 8, member A1
 transforming, acidic coiled-coil containing protein 2
 unti-interacting protein
 procollagen-lysine, 2-oxoglutarate 5-dioxygenase (lysine hydroxylase) 2
 transmembrane 4 superfamily member 7
 triple functional domain (TPTRF) interacting
 hypothetical protein FLJ21343
 small inducible cytokine subfamily C, member 1 (lymphotactin)
 solute carrier family 18 (vesicular monoamine), member 1
 Consensus includes gb:AK024561.1 /DEF=Homo sapiens cDNA: FLJ20508 fls, done ADSE00417, /FEA=mRNA /DB_XREF=gi:10436870 /UG=Hs.306689 Homo sapiens c AK024561
 Consensus includes gb:BC003535.1 /DEF=Homo sapiens, clone IMAGE:3609326, mRNA, partial cds, /FEA=mRNA /PROD=Unknown (protein for IMAGE:3609326) /DB_XI BC003535
 Consensus includes gb:BC003535.1 /DEF=Homo sapiens, clone IMAGE:3609326, mRNA, partial cds, /FEA=mRNA /PROD=Unknown (protein for IMAGE:3609326) /DB_XI BC003535
 Fc fragment of IgE, high affinity I, receptor for; gamma polypeptide
 cysteine and histidine-rich domain (CHORD)-containing, zinc binding protein 1
 chromosome 10 open reading frame 2
 laminin, alpha 4
 G protein coupled receptor interacting protein, complement-c1q tumor necrosis factor-related
 tissue factor pathway inhibitor 2
 nuclear protein, ataxia-telangiectasia locus
 inter-alpha (globulin) inhibitor, H1 polypeptide
 ribosome binding protein 1 homolog 180KD (dog)
 acetyl-Coenzyme A carboxylase alpha
 melanoma antigen, family E, 1, cancer/testis specific
 Consensus includes gb:BC002869.1 /DEF=Homo sapiens, clone IMAGE:3941350, mRNA, partial cds, /FEA=mRNA /PROD=Unknown (protein for IMAGE:3941350) /DB_XI BC002869
 KIAA0508 protein
 G protein-coupled receptor 31
 caspase recruitment domain protein 10
 ubiquitin-conjugating enzyme E2B (RAD6 homolog)
 Consensus includes gb:AF167210 /FEA=EST /DB_XREF=gi:5233719 /DB_XREF=est:tw9405.x1 /CLONE=IMAGE:2400869 /UG=Hs.90449 Human clone 23908 mRNA ser AF167210
 dedicator of cyto-kinesis 2
 epsilon-trimethyllysine hydroxylase
 unc-119 homolog (C. elegans)
 integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41B)
 zinc finger protein 8 (clone HF 18)
 hypothetical protein FLJ23017
 GSC protein
 KIAA0768 protein
 collagen, type XIII, alpha 1
 fibroblast growth factor 7 (keratinocyte growth factor)
 brain expressed, X-linked 1
 hypothetical protein FLJ20294
 calpain 6
 glycoengin 2
 chitinase 3-like 2
 solute carrier family 22 (organic cation transporter), member 1-like antisense
 hypothetical protein FLJ22419
 PR domain containing 1, with ZNF domain
 Splicing factor, arginine/serine-rich, 48KD
 heat shock 90kD protein 1, alpha
 secretagogin
 zinc finger protein 278
 KIAA0476 gene product
 paraoxonase 1
 hypothetical protein FLJ13501
 methylmalonate-semialdehyde dehydrogenase
 KIAA0077 protein
 fibroblast growth factor 18
 Fukuyama type congenital muscular dystrophy
 gb:NM_016407.1 /DEF=Homo sapiens putative integral membrane transporter (LC27), mRNA, /FEA=mRNA /GEN=LC27 /PROD=putative integral membrane transporter /C NM_016407
 peanut-like 1 (Drosophila)

KIAA0551 protein	N25621	1,21
development and differentiation enhancing factor 2	NM_003887	1,21
G protein-coupled receptor 57	NM_014627	1,21
tachykinin receptor 2	NM_001057	1,21
colony stimulating factor 1 receptor, formerly McDonough feline sarcoma viral (v-fms) oncogene homolog	NM_005211	1,21
cyclin-dependent kinase inhibitor 2D (p19, inhibits CDK4)	U20498	1,21
Consensus includes gb:M28219.1 /DEF=Homo sapiens low density lipoprotein receptor (FH 10 mutant causing familial hypercholesterolemia) mRNA, 3 end. /FEA=mRNA /M28219		1,21
polymerase (RNA) II (DNA directed) polypeptide D	BF432147	1,21
NAG16 protein	AF210651	1,21
chondroitin 4-O-sulfotransferase 2	NM_018641	1,21
NAD(P)H dehydrogenase, quinone 1	NM_000903	1,21
solute carrier family 6 (neurotransmitter transporter, creatine), member 8	U17986	1,21
solute carrier family 23 (nucleobase transporters), member 1	AF092511	1,21
Protein inhibitor of activated STAT X	AF077954	1,21
platelet/endothelial cell adhesion molecule (CD31 antigen)	AA702701	1,21
KIAA0451 gene product	AA127643	1,21
transferrin (prealbumin, amyloidosis type I)	AF162690	1,21
diaphanous homolog 2 (Drosophila)	NM_008729	1,21
regulator of G-protein signaling 17	NM_012419	1,21
guanylate cyclase 1, soluble, beta 3	W63728	1,21
prostaglandin D2 synthase (21kD, brain)	BC005939	1,21
G protein-coupled receptor 23	NM_005296	1,21
acetylcholinesterase (YT blood group)	NM_015831	1,21
hypothetical protein FLJ12567	NM_024714	1,21
hypothetical protein FLJ13238	NM_024902	1,21
KIAA1624 protein	NM_020960	1,21
keratin, hair, acidic, 8	NM_006771	1,21
capping protein (actin filament) muscle Z-line, beta	BG420053	1,21
acetyl-Coenzyme A carboxylase alpha	AW188201	1,21
ATPase, Class II, type 9B	RS0094	1,21
IQ motif containing GTPase activating protein 2	NM_006633	1,21
L38424 B subtilis dapB, jpfF, jpfG genes corresponding to nucleotides 1358-3197 of L38424 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	L38424	1,21
very large G protein-coupled receptor 1	AB014586	1,21
reticulon 1	BC000314	1,21
KIAA1535 protein	BF110802	1,21
peter pan homolog (Drosophila)	AI399889	1,21
KIAA0215 gene product	NM_014735	1,21
glial fibrillary acidic protein	NM_002055	1,21
hypothetical protein FLJ10748	NM_018203	1,21
insulin induced gene 1	BG292233	1,21
ribonuclease, RNase A family, 1 (pancreatic)	NM_002933	1,21
zyxin	BC002323	1,21
Consensus includes gb:BE552409 /FEA=EST /DB_XREF=gi:3794101 /DB_XREF=est:hy0708.x1 /CLONE=IMAGE:3196647 /UG=Hs.280895 ESTs		1,21
transmembrane protein with EGF-like and two follistatin-like domains 1	BE552409	1,21
3-hydroxy-3-methylglutaryl-Coenzyme A reductase	BF439316	1,21
succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD	NM_000859	1,21
nuclear receptor subfamily 1, group 1, member 2	NM_003001	1,21
neurogenic differentiation 2	NM_003889	1,21
KIAA1111 protein	AB021742	1,21
cell cycle related kinase	AF091081	1,21
KIAA1660 protein	NM_012119	1,21
transmembrane protein 5	NM_014254	1,21
RCD1 required for cell differentiation1 homolog (S. pombe)	AI360850	1,21
mitogen-activated protein kinase 10	NM_002753	1,20
naturtic peptide receptor A/guanylate cyclase A (atrionatriuretic peptide receptor A)	X15557	1,20
transcription elongation factor B (SII), polypeptide 3 (110kD, elongin A)	NM_003198	1,20
immunoglobulin heavy constant mu	U80164	1,20
pleckstrin and Sec7 domain protein	NM_002779	1,20
histone deacetylase 8	AK024083	1,20
KIAA0672 gene product	NM_014859	1,20
Consensus includes gb:BG032839 /FEA=EST /DB_XREF=gi:12424547 /DB_XREF=est:602300417F1 /CLONE=IMAGE:4401700 /UG=Hs.293341 ESTs, Weakly similar to		1,20
cytochrome P450, subfamily IIIA (nhipedipine oxidase), polypeptide 3	BG032839	1,20
zinc finger protein 281	J04449	1,20
	AK025663	1,20

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Consensus includes gb:BF046750 /FEA=EST /DB_XREF=gi:10763806 /DB_XREF=est:INSSA4 /CLONE=227760 /UG=Hs.274454 ESTs, Weekly similar to 20043994 cthr BF046750
 alpha-methylacyl-CoA racemase
 frizzled homolog 2 (Drosophila)
 LIM domain only 6
 BarH-like homeobox 2
 ketch-like 4 (Drosophila)
 Consensus includes gb:AL133269 /DEF=Human DNA sequence from clone RP3-468K3 on chromosome 8q25-26. Contains a beta tubulin pseudogene, STSs and GSSs /F AL133269
 olfactory receptor, family 7, subfamily E, member 24 pseudogene
 potassium voltage-gated channel, subfamily H (eag-related), member 4
 RAS3D, member RAS oncogene family
 hypothetical protein
 huntingtin-associated protein 1 (neuroan 1)
 alpha-2-HS-glycoprotein
 fibroblast growth factor 2 (basic)
 complement component 8, alpha polypeptide
 cytochrome P450, subfamily IVA, polypeptide 11
 guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1
 cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9
 likely ortholog of preimplantation protein 3
 centaurin, beta 2
 mitogen-activated protein kinase kinase kinase 5
 KIAA0285 gene product
 hypothetical protein FLJ13511
 I-box and WD-40 domain protein 2
 hypothetical protein DKFZ0781P1010
 WAP four-disulfide core domain 1
 KIAA0431 protein
 Consensus includes gb:AA621286 /FEA=EST /DB_XREF=gi:2525225 /DB_XREF=est:z083e09.s1 /CLONE=IMAGE:744616 /UG=Hs.179697 Human DNA damage repair protein
 KIAA0285 gene product
 hypothetical protein IMPACT
 Consensus includes gb:AK025188.1 /DEF=Homo sapiens cDNA: FLJ21535 fts, clone COL08131 /FEA=mRNA /DB_XREF=gi:10437651 /UG=Hs.306781 Homo sapiens cDNA: AK025188
 DKFZP564G202 protein
 estrogen receptor binding site associated, antigen, 9
 hypothetical protein similar to beta-transducin family
 cytochrome P450, subfamily IIA, polypeptide 7
 pre-TNK cell associated protein
 KIAA0747 protein
 DKFZP564G092 protein
 Consensus includes gb:U92027.1 /DEF=Human clone 61501 defective mariner transposon Hsmar2 mRNA sequence, /FEA=mRNA /DB_XREF=gi:2052468 /UG=Hs.15921 U92027
 Consensus includes gb:AV760769 /FEA=EST /DB_XREF=gi:10918617 /DB_XREF=est:AV760769 /CLONE=MDSSE09 /UG=Hs.306324 Homo sapiens mRNA: cDNA DKF AV760769
 protein tyrosine phosphatase, receptor type, K
 hypothetical protein FLJ20920
 hypothetical protein MGC2479
 KIAA0053 gene product
 myelin oligodendrocyte glycoprotein
 gb:NM_023941.1 /DEF=Homo sapiens hypothetical protein MGC3032 (MGC3032), mRNA, /FEA=mRNA /GEN=MGC3032 /PROD=hypothetical protein MGC3032 /DB_XREF=NM_023941
 GW128 protein
 solute carrier family 9 (sodium/hydrogen exchanger), isoform 1 (anionporter, Na+/H+, amiloride sensitive)
 ATPase, H+-transporting, lysosomal (vacuolar proton pump) non-catalytic accessory protein 1B
 KIAA1405 protein
 lysosomal-associated membrane protein 2
 matorin, ring finger protein, 1
 polymerase (DNA directed), delta 3
 aryl hydrocarbon receptor nuclear translocator
 Consensus includes gb:AL137706.1 /DEF=Homo sapiens mRNA: cDNA DKFZP434A2322 (from clone DKFZP434A2322) /FEA=mRNA /PROD=hypothetical protein /DB_XREF=AL137706
 hypothetical protein FLJ20126
 guanine nucleotide binding protein (G protein), alpha 11 (Gq class)
 DNA (cytosine-5-methyltransferase 3-like
 6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 1
 cathepsin B
 EphA7
 RNA binding motif, single stranded interacting protein 1
 Rho GTPase activating protein 1

Consensus includes gb:AB028239 /DEF=Homo sapiens pseudogene for necln /FEA=CDS /DB_XREF=gi:5672590 /UG=Hs.247843 Homo sapiens pseudogene for necln AB028239
 pre-mRNA splicing SR protein RA4 A4648521
 Consensus includes gb:AL049397.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586C1019 (from clone DKFZp586C1019) /FEA=mRNA /DB_XREF=gi:4500188 /UG=Hs.12
 binder of A1 Two BC003087
 KIAA0832 protein AL110250
 potassium voltage-gated channel, shaker-related subfamily, member 1 (episodic ataxia with myokymia) NM_000217
 T cell receptor gamma locus M27331
 KIAA0964 protein BE550798
 KIAA1043 protein AB028966
 hypothetical protein FLJ22028 NM_024854
 androgen-induced prostate proliferative shut-off associated protein NM_015928
 suppression of tumorigenicity 5 NM_005418
 Consensus includes gb:BG494940 /FEA=EST /DB_XREF=gi:13456455 /DB_XREF=IMAGE:4671854 /UG=Hs.554 Sjogren syndrome antigen A BG494940
 Consensus includes gb:BG494940 /FEA=EST /DB_XREF=gi:13456455 /DB_XREF=IMAGE:4671854 /UG=Hs.554 Sjogren syndrome antigen A BG494940
 zinc finger protein 214 NM_000213
 integrin, beta 4 U20760
 calcium-sensing receptor (hypocalcemic hypercalcemia 1, severe neonatal hyperparathyroidism) NM_000293
 Consensus includes gb:AK000293.1 /DEF=Homo sapiens cDNA FLJ20286 fls, clone HEP04358 /FEA=mRNA /DB_XREF=gi:7020278 /UG=Hs.306331 Homo sapiens cDN
 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3 U95430
 hypothetical protein FLJ21019 NM_024927
 endolase alpha, lung-specific NM_014355
 Consensus includes gb:A083904 /FEA=EST /DB_XREF=gi:5811123 /DB_XREF=gi:5811123 /DB_XREF=IMAGE:2561961 /UG=Hs.79732 fibulin 1 A083904
 F-box only protein 24 NM_012172
 hypothetical protein MGC8407 NM_024046
 matrix metalloproteinase 18 (membrane-inserted) U78292
 KIAA0682 gene product NM_014852
 KIAA0329 gene product AB002295
 gb:NM_018069.1 /DEF=Homo sapiens hypothetical protein FLJ10352 (FLJ10352), mRNA /FEA=mRNA /GEN=FLJ10352 /PROD=hypothetical protein FLJ10352 /DB_XREF=gi:5811123 /DB_XREF=IMAGE:2561961 /UG=Hs.79732 fibulin 1 NM_018069
 phosphate cytidylyltransferase 1, choline, alpha isoform AL049542
 DKFZP564F1422 protein NM_024549
 hypothetical protein FLJ21127 NM_004988
 melanoma antigen, family A, 1 (directs expression of antigen MZ2-E) NM_008860
 hydroxyprostaglandin dehydrogenase 15-(NAD) NM_017857
 hypothetical protein FLJ20515 NM_001890
 casein, alpha L06147
 golgi autoantigen, golgin subfamily a, 2 BG169673
 v-rat simian leukemia viral oncogene homolog 8 (ras related; GTP binding protein) BG035151
 splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated) W60800
 hepatic leukemia factor NM_025097
 Consensus includes gb:AK025097.1 /DEF=Homo sapiens cDNA: FLJ21444 fls, clone COL04437 /FEA=mRNA /DB_XREF=gi:10437544 /UG=Hs.306764 Homo sapiens cC
 OLF-1/EBF associated zinc finger gene AW149417
 junctophilin 3 NM_020655
 glycine receptor, alpha 1 (startle disease/hyperkplexia, stiff man syndrome) NM_000171
 integrin, beta 6 NM_000888
 bromodomain adjacent to zinc finger domain, 2A NM_013449
 hypothetical protein FLJ10970 NM_018286
 synovial sarcoma, X breakpoint 3 BF184274
 COX17 homolog, cytochrome c oxidase assembly protein (yeast) NM_005694
 metastasis-associated 1-like 1 AB012922
 C-terminal binding protein 2 AF222711
 retinoic acid receptor, gamma L12060
 a disintegrin and metalloproteinase domain 28 NM_021777
 cryptochrome 1 (photolyase-like) AK022885
 voltage-dependent anion channel 3 D83702
 Consensus includes gb:AK025247.1 /DEF=Homo sapiens cDNA: FLJ21594 fls, clone COL07053 /FEA=mRNA /DB_XREF=gi:10437718 /UG=Hs.288571 Homo sapiens cC
 hypothetical protein FLJ22027 BC002456
 DKFZP564L0862 protein NM_024681
 KIT ligand NM_017757
 interleukin 1 receptor-like 2 NM_024087
 aquaporin 2 (collecting duct) NM_000899
 RNA binding motif protein, Y chromosome, family 1, member A1 NM_003854
 acylserotonin O-methyltransferase-like NM_000466
 NM_005058
 BC002508
 1.19
 1.19

chromosome 1 open reading frame 17
 ATP-binding cassette, sub-family B (MDR/TAP), member 9
 solute carrier family 25 (mitochondrial carrier, Adlar), member 12
 DKFZP38682420 protein
 insulin receptor substrate 3-like
 methylmalonate-semialdehyde dehydrogenase
 chloride intracellular channel 2
 KIAA1418 protein
 Notch homolog 2 (Drosophila)
 arylsulfinase A
 hypothetical protein PRO1048
 thyroid stimulating hormone receptor
 tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator)
 zinc finger protein 142 (clone PHZ-49)
 KIAA0252 protein
 matrix, extracellular phosphoglycoprotein with ASARM motif (bone)
 glycoprotein B (includes Ss blood group)
 glycosylphosphatidylinositol specific phospholipase D1
 ZYG homolog
 RAR-related orphan receptor C
 adaptor-related protein complex 4, epsilon 1 subunit
 phosphoinositide 3-phosphate-binding protein-2
 homeo box B8
 KIAA0144 gene product
 Cdc42 effector protein 2
 Consensus includes gb:AK023058.1 /DEF=Homo sapiens cDNA FLJ12996 fis, clone NTZRP3000235 /FEA=mRNA /DB_XREF=gi:10434801 /UG=Hs.287546 Homo sapiens
 hypothetical protein FLJ14345
 gb:U50073.1 /DEF=Human T cell receptor beta chain precursor (Vbeta 12.3) mRNA, complete cds. /FEA=mRNA /GEN=Vbeta 12.3 /PROD=T cell receptor beta chain precursor
 cadherin 2, type 1, N-cadherin (neuronal)
 Consensus includes gb:AW517464 /FEA=EST /DB_XREF=gi:7155546 /DB_XREF=ast:xp94g06.x1 /CLONE=IMAGE:2748058 /UG=Hs.119598 ribosomal protein L3
 hypothetical protein FLJ22222
 casein kinase 1, epsilon
 myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 10
 dihydroorotate dehydrogenase
 palladin
 hypothetical protein FLJ13373
 ELK3, ETS-domain protein (SRF accessory protein 2)
 interleukin 18 receptor 1
 bone morphogenetic protein 7 (osteogenic protein 1)
 lens intrinsic membrane protein 2 (19KD)
 downregulated in ovarian cancer 1
 squamous cell carcinoma antigen recognised by T cells 3
 interleukin 2 receptor, alpha
 KIAA0387 protein
 fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome)
 alkaline phosphatase, placental-like 2
 microphthalmia-associated transcription factor
 dishevelled, dish homolog 2 (Drosophila)
 mucin 3A, intestinal
 PRO1748 protein
 hypothetical protein FLJ10636
 deleted in lung and esophageal cancer 1
 protocadherin alpha 2
 neuronal PAS domain protein 2
 orosomucoid 1
 hypothetical protein FLJ00001
 Consensus includes gb:AU146050 /FEA=EST /DB_XREF=gi:1007571 /DB_XREF=ast:AU146050 /CLONE=HEMBA1006665 /UG=Hs.287459 Homo sapiens cDNA FLJ11111
 Tax1 (human T-cell leukemia virus type I) binding protein 1
 KIAA0125 gene product
 proopiomelanocortin (adrenocorticotropin/ beta-lipotropin/ alpha-melanocyte stimulating hormone/ beta-melanocyte stimulating hormone/ beta-endorphin)
 immune dysregulation, polyendocrinopathy, enteropathy, X-linked
 beta-1,3-glucuronidyltransferase 1 (glucuronosyltransferase P)
 Prader-Willi/Angelman syndrome-5

protein kinase NYD-SP15
 KIA0583 gene product
 PRO1578 protein
 heparan sulfate (glucosamine) 3-O-sulfotransferase 1
 hypothetical protein FLJ23053
 CGI-101 protein
 Consensus includes gb:AF131844.1 /DEF=Homo sapiens clone 25015 mRNA sequence. /FEA=mRNA /DB_XREF=gi:4406685 /UG=Hs.201626 Homo sapiens clone 25015 AF131844
 Consensus includes gb:AK024457.1 /DEF=Homo sapiens mRNA for FLJ00049 protein, partial cds. /FEA=mRNA /GEN=FLJ00049 /PROD=FLJ00049 protein /DB_XREF=gi:AK024457
 mitogen-activated protein kinase kinase 3
 leucine-rich repeat protein, neuronal 3
 ret proto-oncogene (multiple endocrine neoplasia and medullary thyroid carcinoma 1, Hirschsprung disease)
 frequenin homolog (Drosophila)
 Kruppel-like factor 12
 transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)
 Notch homolog 2 (Drosophila)
 pregnancy specific beta-1-glycoprotein 1
 H4 histone family, member J
 retinol binding protein 4, plasma
 jerky homolog (mouse)
 glutamate receptor, metabotropic 7
 phospholamban
 sulfotransferase family, cytosolic, 2A, dehydroepiandrosterone (DHEA)-preferring, member 1
 sirinuin silent mating type information regulation 2 homolog 2 (S. cerevisiae)
 forkhead box D1
 tropomyosin T3, skeletal, fast
 myeloid/lymphoid or mixed-lineage leukemia (littorax homolog, Drosophila)
 Consensus includes gb:AK026188.1 /DEF=Homo sapiens cDNA: FLJ22535 fls, clone HRC13115, highly similar to AF152336 Homo sapiens protocadherin gamma B7 (PCI AK026188)
 hypothetical protein PP1345
 KIAA1075 protein
 chondromodulin 1 precursor
 Consensus includes gb:AF070575.1 /DEF=Homo sapiens clone 24407 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3387946 /UG=Hs.12432 Homo sapiens clone 24407 AF070575
 T cell receptor alpha locus
 plakophilin 4
 nuclear receptor co-repressor 1
 paired basic amino acid cleaving system 4
 Rhesus blood group, D antigen
 tissue inhibitor of metalloproteinase 3 (Sorby fundus dystrophy, pseudoinflammatory)
 KIAA1609 protein
 hypothetical protein FLJ13285
 Consensus includes gb:A523613 /FEA=EST /DB_XREF=gi:4437748 /DB_XREF=est:ig95a07.x1 /CLONE=IMAGE:2118500 /UG=Hs.293495 ESTs, Weakly similar to ALU1
 pre-mRNA processing factor 18
 KIAA0863 protein
 myosin VI
 Consensus includes gb:l42230 /DEF=Homo sapiens core 2 beta-1,6-N-acetylglucosaminyltransferase pseudogene /FEA=CDS /DB_XREF=gi:1018163 /UG=Hs.247990 Ho L42230
 Consensus includes gb:AU148957 /FEA=EST /DB_XREF=gi:11010478 /DB_XREF=est:AU148957 /CLONE=NT2RM4001313 /UG=Hs.306638 Homo sapiens cDNA FLJ12: AU1543
 chromodomain helicase DNA binding protein 3
 small inducible cytokine subfamily A (Cys-Cys), member 18, pulmonary and activation-regulated
 hypothetical protein FLJ10572
 hypothetical protein FLJ14033 similar to hypoxia inducible factor 3, alpha subunit
 microseminoprotein, beta-
 synrophin, beta 2 (dystrophin-associated protein A1, 59KD, basic component 2)
 PRY2 N-terminal domain-interacting receptor 1
 acid phosphatase, prostate
 small inducible cytokine subfamily D (Cys-X3-Cys), member 1 (fractalkine, neurotactin)
 RAW binding protein 1
 integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)
 hypothetical protein FLJ20731
 uronyl 2-sulfotransferase
 ubiquitin-conjugating enzyme E2A (RAD6 homolog)
 hypothetical protein DKFZp434G2228
 DKFZP434K081 protein
 hypothetical protein FLJ20574
 integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)

tryptase beta 1
 Consensus includes gb:AK024909.1 /DEF=Homo sapiens cDNA: FLJ21258 f1, clone COL01402. /FEA=mRNA /DB_XREF=gi:10437324 /UG=Hs.306724 Homo sapiens cDNA: AK024909
 hypothetical protein FLJ12178
 peptide YY
 retinoic acid receptor, alpha
 CD3E antigen, epsilon polypeptide (TIT3 complex)
 Consensus includes gb:BE868287 /FEA=EST /DB_XREF=gi:10340242 /DB_XREF=est:60150987 /F1 /CLONE=IMAGE:3911808 /UG=Hs.249219 Homo sapiens chromosome BE868287
 interleukin 18 receptor accessory protein
 hypothetical protein FLJ21120
 cyclin, basic protein of sperm head cytoskeleton 1
 zinc-finger protein DZP1
 neurotrophin FF-amide peptide precursor
 adrenergic, beta, receptor kinase 2
 Consensus includes gb:AL353949.1 /DEF=Homo sapiens mRNA: cDNA DKFp761P1114 (from clone DKFp761P1114). /FEA=mRNA /DB_XREF=gi:7669990 /UG=Hs.301
 telomeric repeat binding factor (NIMA-interacting) 1
 parvin, alpha
 small proline-rich protein 2C
 proboscoid gamma subfamily A, 3
 much 4, tracheobronchial
 nucleoporin 50K0
 serine/threonine kinase 38 like
 hypothetical protein DKFp566H0824
 solute carrier family 7 (cationic amino acid transporter, y+ system), member 9
 zinc finger protein 237
 growth differentiation factor 9
 hypothetical protein KIAA0244 protein
 putative dipeptidase
 protein phosphatase 3 (formerly 2B), catalytic subunit, gamma isoform (calcineurin A gamma)
 sphingomyelin phosphodiesterase 2, neutral membrane (neutral sphingomyelinase)
 STAT induced STAT inhibitor-2
 hypothetical protein FLJ22662
 adrenomedullin receptor
 mairlin 2
 KIAA0459 protein
 PRO1900 protein
 bombesin-like receptor 3
 synovial sarcoma, X breakpoint 2
 Burkitt lymphoma receptor 1, GTP binding protein
 synaptobegmin 1
 collagen, type VI, alpha 1
 reticulon 1
 hypothetical protein FLJ14529
 sh3-associated polypeptide, 30K0
 nuclear RNA export factor 2
 nuclear transcription factor Y, beta
 Consensus includes gb:AW574818 /FEA=EST /DB_XREF=gi:7246357 /DB_XREF=est:UI-HF-BK0-abi-e-06-0-UI.s1 /CLONE=IMAGE:3056362 /UG=Hs.32374 ESTs, Weak
 venth 1
 phospholipase D2
 DKFZP586D0919 protein
 KIAA0871 protein
 arnedillo repeat gene deletions in velocardiofacial syndrome
 Consensus includes gb:BF509345 /FEA=EST /DB_XREF=gi:11592709 /DB_XREF=est:UI-H-B14-aox-c-11-0-UI.s1 /CLONE=IMAGE:3086229 /UG=Hs.6996 ESTs
 glutathione S-transferase M4
 signal recognition particle receptor (docking protein)
 hypothetical protein FLJ12586
 coronin, actin binding protein, 2A
 phenylalanine-tRNA synthetase
 Lem3 protein
 PTK2 protein tyrosine kinase 2
 calcium channel, voltage-dependent, beta 2 subunit
 oxidised low density lipoprotein (lectin-like) receptor 1
 Consensus includes gb:AU144791 /FEA=EST /DB_XREF=gi:11006312 /DB_XREF=est:AU144791 /CLONE=HEMBA1002971 /UG=Hs.274237 Homo sapiens cDNA AU144791

smoothed homolog (Drosophila) NM_005631 1.19
small inducible cytokine A4 NM_002984 1.19
insulin-like growth factor 2 (somatomedin A) NM_000612 1.19
Consensus includes gb:AF052172.1 /DEF=Homo sapiens clone 24617.1 /AF052172 1.19
protein (epitope-prolyl dipeptidyl aminopeptidase) NIMA-interacting, 4 (osculin) NM_008223 1.19
Consensus includes gb:AF070157.1 /FEA=EST /DB_XREF=gi:5766983 /DB_XREF=est:wn89h08.x1 /CLONE=IMAGE:2479263 /UG=Hs.77631 glycine cleavage system protei A1970157 1.19
hypothetical protein FLJ12606 NM_024604 1.19
Consensus includes gb:AW138468 /FEA=EST /DB_XREF=gi:6142786 /DB_XREF=est:UH-B1-adtg-401-Q-U1.s1 /CLONE=IMAGE:2716920 /UG=Hs.299882 ESTs, Highly: AW138468 1.19
Consensus includes gb:AF090887.1 /DEF=Homo sapiens clone HQ0085 /FEA=mrna /DB_XREF=gi:6690154 /UG=Hs.306562 Homo sapiens clone HQ0085 1.19
spermatogenesis associated 2 NM_006038 1.19
gamma-aminobutyric acid (GABA) B receptor, 1 AL031883 1.19
Purkinje cell protein 4 NM_006198 1.19
transforming growth factor, beta 2 M16154 1.19
hypothetical protein FLJ20014 NM_017622 1.19
glycoprotein hormones, alpha polypeptide NM_000735 1.19
elastin (supraaortic stenosis, Williams-Beuren syndrome) M24782 1.19
Consensus includes gb:AL121964 /DEF=Human DNA sequence from clone RP1-154G14 on chromosome 8q15-16.3. Contains the 3' end of the MAP3K7 gene for mitogen- AF212895 1.19
cullin 4B NM_002747 1.19
mitogen-activated protein kinase 4 AF023450 1.19
Down syndrome cell adhesion molecule NM_006942 1.19
SRY (sex determining region Y)-box 20 NM_012424 1.19
ribosomal protein S6 kinase, 52kD, polypeptide 1 NM_001709 1.19
brain-derived neurotrophic factor NM_021911 1.19
gamma-aminobutyric acid (GABA) A receptor, beta 2 NM_012390 1.19
protein homologous to salivary proline-rich protein P-8 NM_001508 1.19
G protein-coupled receptor 39 NM_0089513 1.19
Consensus includes gb:AF089513 /FEA=EST /DB_XREF=gi:5594677 /DB_XREF=est:wn05e06.x1 /CLONE=IMAGE:2444578 /UG=Hs.168694 Homo sapiens clone 23763 u A089513 1.19
microtubule-associated protein, RPIIB family, member 3 BG222594 1.19
glucosylceramide/CO77 synthase; Gb3/CO77 synthase; alpha1,4-galactosyltransferase; 4-N-acetylglucosaminyltransferase NM_017436 1.19
NEU1 protein NM_016215 1.19
hypothetical protein BF787555 1.19
non-canonical ubiquitin conjugating enzyme 1 AW500009 1.19
SECT homolog NM_012455 1.19
heat shock 27kD protein family, member 7 (cardiovascular) NM_014424 1.19
Consensus includes gb:AK024879.1 /DEF=Homo sapiens cDNA: FLJ12228 fls. done COL00721. /FEA=mrna /DB_XREF=gi:10437291 /UG=Hs.308715 Homo sapiens cDNA: AK024879 1.19
putative protein-tyrosine kinase NM_015978 1.19
Consensus includes gb:AF056433.1 /DEF=Homo sapiens clone FBD3 Cn-du-chat critical region mRNA. /FEA=mrna /DB_XREF=gi:3044151 /UG=Hs.8078 Homo sapiens AF056433 1.19
hypothetical protein MGC5350 NM_030920 1.19
zinc finger protein 26 (KOX 20) NM_019591 1.19
multiple endocrine neoplasia 1 NM_000244 1.19
protein-L-isoaspartate (D-aspartate) O-methyltransferase NM_005389 1.19
M11167 Human 28S rRNA sequence, length 5025 bases, 5 prime target bases 1-1665 M11167 1.19
dihydropyrimidinase-like 3 W72516 1.19
Consensus includes gb:AK022309.1 /DEF=Homo sapiens cDNA FLJ12247 fls. clone MAMMA1001397. /FEA=mrna /DB_XREF=gi:10433678 /UG=Hs.287506 Homo sapiens AK022309 1.19
N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 2 NM_003635 1.19
Consensus includes gb:AA102667 /FEA=EST /DB_XREF=gi:1647984 /DB_XREF=est:zn73e05.s1 /CLONE=IMAGE:563840 /UG=Hs.294673 ESTs, Weakly similar to ALU8 AA102667 1.19
glypican 4 NM_001448 1.19
hypothetical protein NM_019103 1.19
adaptor-related protein complex 3, sigma 2 subunit BC002785 1.19
Consensus includes gb:BE046461 /FEA=EST /DB_XREF=gi:8363514 /DB_XREF=est:hn47g05.x2 /CLONE=IMAGE:3026840 /UG=Hs.159412 Homo sapiens clone 24628 r BE046461 1.19
hypothetical protein FLJ20075 NM_017655 1.19
fasciculation and elongation protein zeta 1 (zyglin I) NM_022549 1.19
cytochrome P450, subfamily 11C (mephenytoin 4-hydroxylase), polypeptide 8 NM_030878 1.19
G protein-coupled receptor 9 NM_001504 1.19
hypothetical protein FLJ20005 NM_017817 1.19
hypothetical protein FLJ10865 NM_018173 1.19
RAS11B, member RAS oncogene family X79780 1.19
mitogen-activated protein kinase kinase 13 NM_004721 1.19
dynen, axonemal, intermediate polypeptide, 1 NM_012144 1.19
carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein) M69176 1.19
troponin T2, cardiac X79857 1.19
tumor suppressing subtransferable candidate 3 BC005034 1.19
calmeglin NM_004362 1.19

potassium inwardly-rectifying channel, subfamily J, member 3
hypothetical protein FLJ20958
paired box gene 4
Consensus includes gb:AL520875 /FEA=EST /DB_XREF=gi:12784168 /DB_XREF=est:AL520875 /CLONE=CS0DB002YF15 (3 prime) /UG=Hs.77783 membrane-associate
Consensus includes gb:AF141801.1 /DEF=Homo sapiens GKAPSAPAP interacting protein (SHANK) mRNA, partial cds. /FEA=mRNA /GEN=SHANK /PROD=GKAPSAPAF AF141801
hypothetical protein FLJ11186
platelet factor 4 variant 1
senna domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3F
ribosome binding protein 1 homolog 180kD (dog)
sperm protein associated with the nucleus, X chromosome, family member A1
hypothetical protein FLJ11183
alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, S. cerevisiae)
claudin 14
alcohol dehydrogenase 1A (class I), alpha polypeptide
cell division cycle 25B
hypothetical protein F23149_1
fucosyltransferase 3 (Galactoside 3(4)-L-fucosyltransferase, Lewis blood group included)
Consensus includes gb:AA417256 /FEA=EST /DB_XREF=gi:2077373 /DB_XREF=est:zu07e04.s1 /CLONE=IMAGE:731166 /UG=Hs.193769 ESTs
KIA00349 protein
hypothetical protein FLJ10292
cleavage stimulation factor, 3' pre-RNA, subunit 1, 50kD
guanine nucleotide binding protein-like 1
integral inner nuclear membrane protein
sperm associated antigen 6
prostaglandin E receptor 1 (subtype EP1), 42kD
Consensus includes gb:AV700298 /FEA=EST /DB_XREF=gi:10302269 /DB_XREF=est:AV700298 /CLONE=GKCBVG05 /UG=Hs.285173 ESTs, Moderately similar to CD4 AV700298
nucleoporin 98kD
Consensus includes gb:AF070810.1 /DEF=Homo sapiens clone 24505 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3387982 /UG=Hs.100543 Homo sapiens clone 24505 AF070810
hypothetical protein FLJ10634
hypothetical protein MGC10763
suppressor of S. cerevisiae gcr2
nuclear receptor subfamily 4, group A, member 1
neuronal pentraxin II
Consensus includes gb:AU154785 /FEA=EST /DB_XREF=gi:1016306 /DB_XREF=est:AU154785 /CLONE=NT2RPA002888 /UG=Hs.31532 Homo sapiens mRNA; cDNA (AU154785
RAD52 homolog (S. cerevisiae)
hypothetical protein DKFZP586M1120
DKFZP586B0923 protein
profilin 2
serine protease inhibitor, Kazal type 2 (acrosin-tyrosin inhibitor)
centrin, EF-hand protein, 3 (CDC31 homolog, yeast)
gamma-aminobutyric acid (GABA) A receptor, alpha 4
Consensus includes gb:BC000975.2 /DEF=Homo sapiens, clone IMAGE:3448306, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3448306) /DB_XI BC000975
CD84 antigen (leukocyte antigen)
hypothetical protein FLJ22757
hypothetical protein FLJ11729
DKFZP566A0946 protein
actinin, alpha 1
calpain 2, (mu) large subunit
Fc fragment of IgA, receptor for
SEC24 related gene family, member B (S. cerevisiae)
protein tyrosine phosphatase, non-receptor type 9
solute carrier family 12 (sodium/chloride transporters), member 3
alkylation repair, alk8 homolog
necroline receptor 2 (cardiac)
small inducible cytokine subfamily A (Cys-Cys), member 13
ataxia telangiectasia mutated (includes complementation groups A, C and D)
clone FLB5214
intercellular adhesion molecule 5, telencephalin
solute carrier family 7, cationic amino acid transporter, y+ system) member 10
Consensus includes gb:AK027217.1 /DEF=Homo sapiens cDNA: FLJ23564 fls, clone LNG10773, /FEA=mRNA /DB_XREF=gi:10440296 /UG=Hs.306913 Homo sapiens cDNA: AK027217
transglutaminase 5
hypothetical protein FLJ20507
nucleolar protein family A, member 2 (HACA small nucleolar RNPs)

nuclear receptor coactivator 3	NM_006534	1.18
Consensus includes gb:AK025177.1 /DEF=Homo sapiens cDNA: FLJ21524 fts, clone COL05921 /FEA=mRNA /DB_XREF=gi:10437640 /UG=Hs.306778 Homo sapiens cDNA: AK025177	NM_000116	1.18
tariazin (cardiomyopathy, dilated 3A (X-linked); endocardial fibroelastosis 2; Barth syndrome)	NM_004375	1.18
COX11 homolog, cytochrome c oxidase assembly protein (yeast)	NM_000900	1.18
matrix Gla protein	NM_024732	1.18
hypothetical protein FLJ14351	NM_004670	1.18
3'-phosphoadenosine 5'-phosphosulfate synthase 2	NM_005832	1.18
potassium large conductance calcium-activated channel, subfamily M, beta member 2	NM_005399	1.18
protein kinase, AMP-activated, beta 2 non-catalytic subunit	AV700403	1.18
Consensus includes gb:AV700403 /FEA=EST /DB_XREF=gi:10302374 /DB_XREF=est:AV700403 /CLONE=GK6ADA01 /UG=Hs.293673 ESTs	NM_003201	1.18
transcription factor A, mitochondrial	A1246541	1.18
hypothetical protein FLJ14154	NM_025163	1.18
hypothetical protein FLJ12768	BF509069	1.18
hypothetical protein MGC202727	AL109874	1.18
Consensus includes gb:AL109874.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 46866 /FEA=mRNA /DB_XREF=gi:5688832 /UG=Hs.221811	NM_020250	1.18
MOST2 protein	NM_024706	1.18
hypothetical protein FLJ13479	NM_022912	1.18
hypothetical protein FLJ13110	NM_003645	1.18
fatty-acid-Coenzyme A ligase, very long-chain 1	NM_002935	1.18
ribonuclease, RNase A family, 3 (eosinophil cationic protein)	AF265167	1.18
ATP-binding cassette, sub-family A (ABC1), member 1	AF239764	1.18
EGF-like module-containing mucin-like receptor EMR3	NM_000810	1.18
gamma-aminobutyric acid (GABA) A receptor, alpha 5	NM_015880	1.18
RIG-like 14-1	NM_021235	1.18
epidermal growth factor receptor substrate EPS15R	A1358867	1.18
apolipoprotein E	A1837543	1.18
DC12 protein	NM_001981	1.18
epidermal growth factor receptor pathway substrate 15	AC004755	1.18
one out domain, family member 3	A1963043	1.18
Consensus includes gb:A1963043 /FEA=EST /DB_XREF=gi:5810262 /DB_XREF=est:hwz30b11.x1 /CLONE=IMAGE:2359549 /UG=Hs.246885 hypothetical protein FLJ2078	AV651241	1.18
Consensus includes gb:AL049242.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564B083 (from clone DKFZp564B083) /FEA=mRNA /DB_XREF=gi:4499977 /UG=Hs.2347	AA046411	1.18
Consensus includes gb:AV651241 /FEA=EST /DB_XREF=gi:9872255 /DB_XREF=est:AV651241 /CLONE=GLCCNG03 /UG=Hs.282469 ESTs	AF188773	1.18
amyloid beta precursor protein (cytoplasmic tail) binding protein 2	NM_004636	1.18
dual-specific tyrosine-(Y)-phosphorylation regulated kinase 3	U66584	1.18
sema domain, immunoglobulin domain (lg), short basic domain, secreted, (semaphorin) 3B	AF109294	1.18
crystallin, alpha A	AK022102	1.18
methylnucleoside phosphorylase	NM_008071	1.18
Consensus includes gb:AK022102.1 /DEF=Homo sapiens cDNA FLJ12040 fts, clone HEMBB1001944 /FEA=mRNA /DB_XREF=gi:10433423 /UG=Hs.296687 Homo sapiens cDNA: AK022102	NM_005380	1.18
polycystic kidney disease (polycystin) and REJ (sperm receptor for egg jelly, sea urchin homolog)/like	A1796813	1.18
v-maf musculoaponeurotic fibrosarcoma oncogene homolog (avian)	NM_005048	1.18
adipilin	AL046684	1.18
parathyroid hormone receptor 2	L00692	1.18
retinoblastoma (X-linked, juvenile) 1	AF007141	1.18
carcinoembryonic antigen-related cell adhesion molecule 3	AW268840	1.18
5-hydroxytryptamine (serotonin) receptor 6	NM_005756	1.18
eukaryotic translation initiation factor 4E	U94506	1.18
G protein-coupled receptor 64	AK023069	1.18
tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)	AL574660	1.18
hypothetical protein	NM_022820	1.18
ATP-binding cassette, sub-family D (ALD), member 4	AW026846	1.18
cytochrome P450 polypeptide 43	NM_002939	1.18
C1q-related factor	A1023317	1.18
ribonuclease/angiogenin inhibitor	NM_024582	1.18
KIAA0130 gene product	NM_023922	1.18
hypothetical protein FLJ23056	NM_006355	1.18
taste receptor, type 2, member 14	A1341811	1.18
ring finger protein 15	NM_001386	1.18
gene amplified in squamous cell carcinoma 1	X88405	1.18
dihydropyrimidinase-like 2	U57001	1.18
myelin associated glycoprotein	NM_006890	1.18
ephrin-B3	NM_002585	1.18
carcinoembryonic antigen-related cell adhesion molecule 7		
pre-B-cell leukemia transcription factor 1		

Consensus includes gb:AK023816.1 /DEF=Homo sapiens cDNA FLJ13754 fls. clone PLACE3000362. /FEA=mRNA /DB_XREF=gj:10435965 /UG=Hs.136233 Homo sapiens inhibin, alpha
 1.18 MI3881

NIMA (never in mitosis gene a)-related kinase 1
 1.18 AF155113

peroxisome proliferative activated receptor, alpha
 1.18 NM_005036

MAD, mothers against decapentaplegic homolog 6 (Drosophila)
 1.18 AF035528

hypothetical protein PP1628
 1.18 NM_025201

hypothetical protein DKFZ434C135
 1.18 NM_030938

ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome
 1.18 AF000895

lucosyltransferase 6 (alpha (1.3) lucosyltransferase)
 1.18 U27335

cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 6
 1.18 AF182275

AD037 protein
 1.18 AI890181

FXYD domain-containing ion transport regulator 7
 1.18 NM_022006

KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3
 1.18 NM_016657

MHC class I region ORF
 1.18 NM_006674

TGFB inducible early growth response 2
 1.18 AA146594

hypothetical protein FLJ23259
 1.18 NM_024727

J domain containing protein 1
 1.18 NM_021800

sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F
 1.18 NM_004263

angiotensin receptor-like 2
 1.18 NM_005162

sperm protein associated with the nucleus, X chromosome, family member A1
 1.18 NM_013453

Consensus includes gb:BF344237 /FEA=EST /DB_XREF=gj:1291450 /DB_XREF=est:60201749 /f1 /CLONE=IMAGE:4152983 /UG=Hs.124776 Homo sapiens mRNA, cc BF344237
 1.18 BE964082

complement component 1, q subcomponent binding protein
 1.18 NM_005312

guanine nucleotide-releasing factor 2 (specific for crk proto-oncogene)
 1.18 AI816291

hypothetical protein
 1.18 BF440025

nephroblastoma overexpressed gene
 1.18 gb:AF165520.1 /DEF=Homo sapiens phorbol 1 protein (PBI) mRNA, complete cds. /FEA=mRNA /GEN=PBI /PROD=phorbol 1 protein /DB_XREF=gj:9294746 /UG=Hs.85 AF165520

astrotactin
 1.18 AB006627

ATPase, Class I, type 8B, member 2
 1.18 AL137537

hypothetical protein MGC4659
 1.18 NM_025268

collagen, type V, alpha 2
 1.18 AL575735

interleukin 4
 1.18 NM_000589

helicase-mot
 1.18 AK001827

histamine receptor H1
 1.18 D28481

Consensus includes gb:BE965029 /FEA=EST /DB_XREF=gj:11768976 /DB_XREF=est:601658812R1 /CLONE=IMAGE:3886131 /UG=Hs.198793 Homo sapiens cDNA, FL BE965029
 1.18 AB051441

KIAA1654 protein
 1.18 NM_002392

Mdm2, transformed 3T3 cell double minute 2, p53 binding protein (mouse)
 1.18 L17328

fasciculation and elongation protein zeta 2 (zyglin II)
 1.18 AI424243

CEGP1 protein
 1.18 NM_003641

Interferon induced transmembrane protein 1 (9-27)
 1.18 NM_022898

B-cell CLL/lymphoma 11B (zinc finger protein)
 1.18 AK024051

MUF1 protein
 1.18 NM_021975

v-rel reticuloendotheliosis viral oncogene homolog A, nuclear factor of kappa light polypeptide gene enhancer in B-cells 3, p65 (avian)
 1.18 BC008245

fibroblast growth factor 18
 1.18 AF004231

leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2
 1.18 BC002665

proteolipid protein 1 (Pelizaeus-Merzbacher disease, spastic paraplegia 2, uncomplicated)
 1.18 AL045513

protein O-lucosyltransferase 1
 1.18 AB036432

advanced glycosylation end product-specific receptor
 1.18 NM_005751

A kinase (PKA) anchor protein (yoliao) 9
 1.18 AL050308

NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3 (12kD, B12) pseudogene 5
 1.18 BC003154

tripartite motif-containing 32
 1.18 AF081194

ectodermal dysplasia 1, anhidrotic
 1.18 BC001362

nucleotide-sugar transporter similar to C. elegans sqv-7
 1.18 AJ005866

2',3'-cyclic nucleotide 3' phosphodiesterase
 1.18 NM_014374

gb:U36759.1 /DEF=Human pre-T cell receptor alpha-type chain precursor, mRNA, complete cds. /FEA=mRNA /PROD=pre-T cell receptor alpha-type chain precursor /DB_XREF=gj:10435965 /UG=Hs.136233 Homo sapiens pre-T cell receptor alpha-type chain precursor
 1.18 U36759

replication initiation region protein (60kD)
 1.18 BE973687

hair cell homology (Drosophila)
 1.18 NM_022647

phosphoinositide-3-kinase, class 3
 1.18 AF263310

rab3 interacting protein
 1.18 NM_016158

erythrocyte transmembrane protein
 1.18 AI359472

KIAA1098 protein
 1.18 AA515698

tubulin, beta, 2
 1.18 X88405

myelin associated glycoprotein
 1.18 NM_005235

v-erb-a erythroblastic leukemia viral oncogene homolog 4 (avian)

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Consensus includes gb:AL442084.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547G044 (from clone DKFZp547G044). /FEA=mRNA /DB_XREF=gi:10241763 /UG=Hs.306 AL442084
 G protein-coupled receptor 52 NM_005684 1.18
 KIA00750 gene product NM_014632 1.18
 ATP-binding cassette, sub-family G (WHITE), member 2 AF098951 1.18
 ras responsive element binding protein 1 AW118862 1.18
 Consensus includes gb:AK025086.1 /DEF=Homo sapiens cDNA: FLJ21433 fs, clone COL04232. /FEA=mRNA /DB_XREF=gi:10437530 /UG=Hs.306761 Homo sapiens cDNA: AK025086
 homeo box C13 NM_017410 1.18
 HYA22 protein BF031714 1.18
 calcium channel, voltage-dependent, beta 2 subunit NM_000724 1.18
 adhesion glycoprotein NM_005666 1.18
 H2B histone family, member D NM_003520 1.18
 adenomatous polyposis coli NM_000038 1.18
 a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 12 NM_030955 1.18
 adrenergic, alpha-2A, receptor AF284095 1.18
 calcium binding protein 2 NM_016366 1.18
 amphiregulin (schwannoma-derived growth factor) NM_001657 1.18
 annexin A8 NM_001630 1.18
 H1 histone family, member 4 AL353759 1.18
 ribonucleotide reductase M1 polypeptide NM_001033 1.18
 Consensus includes gb:AI590869 /FEA=EST /DB_XREF=gi:4599517 /DB_XREF=IMAGE:2266804 /UG=Hs.172035 hypothetical protein similar to AI590869
 H3 histone, family 3A A1955855 1.18
 indoleamine-pyrole 2,3 dioxygenase M34455 1.18
 Consensus includes gb:AL137113.1 /DEF=Homo sapiens mRNA; cDNA DKFZp4340523 (from clone DKFZp4340523). /FEA=mRNA /DB_XREF=gi:6808127 /UG=Hs.3064 AL137113
 KIA00887 protein AB020694 1.18
 period homolog 1 (Drosophila) NM_002616 1.18
 mab-21-like 2 (C. elegans) AF262032 1.18
 heat shock 70kD protein 8 (HSP70B) NM_002155 1.18
 hypothetical protein FLJ11040 BF668108 1.18
 forkhead box Q3A AF041336 1.18
 Consensus includes gb:AL049313.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564B076 (from clone DKFZp564B076). /FEA=mRNA /DB_XREF=gi:4500086 /UG=Hs.2110: AL049313
 Consensus includes gb:BG149837 /FEA=EST /DB_XREF=gi:12661867 /DB_XREF=IMAGE:3434062 /UG=Hs.33540 ESTs, Weakly similar to dL: BG149837
 hepatocellular carcinoma antigen gene 520 NM_022087 1.18
 hypothetical protein CDA08 NM_030790 1.18
 MyoD family inhibitor NM_005586 1.18
 Consensus includes gb:LA8728 /DEF=Homo sapiens T cell receptor beta (TCRBV10S1) gene, complete cds /FEA=mRNA_1 /DB_XREF=gi:1054550 /UG=Hs.37163 Homo LA8728 1.18
 Testis-specific XK-related protein on Y NM_004677 1.18
 gb:Z25428.1 /DEF=H. sapiens protein-serine/threonine kinase gene, complete cds. /FEA=mRNA /PRODB=protein-serine/threonine kinase /DB_XREF=gi:405734 /FL=gb:Z25428
 KIA00850 protein AB014550 1.18
 myosin IXA NM_006901 1.18
 KIA00800 gene product NM_014703 1.18
 hypothetical protein FLJ10210 NM_018027 1.18
 small inducible cytokine B subfamily (Cys-X-Cys motif), member 13 (B-cell chemoattractant) NM_006419 1.17
 small proline-rich protein 18 (cornalin) NM_003125 1.17
 surfactant, pulmonary-associated protein C N56912 1.17
 alpha thalassemia/mental retardation syndrome X-linked (RAD54 homolog, S. cerevisiae) BC002521 1.17
 fetuin B AB017551 1.17
 Consensus includes gb:AL137428.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761N1323 (from clone DKFZp761N1323). /FEA=mRNA /DB_XREF=gi:6807885 /UG=Hs.30 AL137428 1.17
 Consensus includes gb:AL049888.1 /DEF=Homo sapiens mRNA; cDNA DKFZp584F212 (from clone DKFZp584F212). /FEA=mRNA /DB_XREF=gi:4884239 /UG=Hs.3063 AL049888 1.17
 interleukin 11 receptor, alpha NM_004512 1.17
 claudin 5 (transmembrane protein deleted in velocardiofacial syndrome) NM_003277 1.17
 Consensus includes gb:AL117652.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586B1324 (from clone DKFZp586B1324). /FEA=mRNA /DB_XREF=gi:5912244 /UG=Hs.271 AL117652 1.17
 pan-hematopoietic expression NM_005705 1.17
 angiotensin I converting enzyme (peptidyl-dipeptidase A) 2 NM_021804 1.17
 KIA00321 protein AB002319 1.17
 chromosome 6 open reading frame 5 AW296296 1.17
 PR domain containing 13 NM_021620 1.17
 basic leucine zipper nuclear factor 1 (JEM-1) AF286161 1.17
 Huntingtin interacting protein H AF161412 1.17
 KIA00323 protein AU075450 1.17
 ATP-binding cassette, sub-family D (ALD), member 2 NM_005164 1.17
 KIA00471 gene product AB019480 1.17
 T-cell leukemia/lymphoma 6 NM_020552 1.17
 hypothetical protein FLJ20909 NM_024693 1.17

hypothetical gene CO030
 inducible T-cell co-stimulator
 protein kinase (cAMP-dependent, catalytic) inhibitor gamma
 CGI-121 protein
 Consensus includes gp:AL049992.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G242 (from clone DKFZp564G242). /FEA=mRNA /DB_XREF=gi:4884244 /UG=Hs.2258 /AL049992
 L47125
 glypican 3
 hypothetical protein MGC3146
 myosin IXB
 hepatocyte nuclear factor 3, alpha
 matrix metalloproteinase 11 (stromelysin 3)
 keratin 12 (Meesmann corneal dystrophy)
 crystallin, gamma B
 dipeptidylpeptidase IV (CD26, adenosine deaminase complementing protein 2)
 X-ray repair complementing defective repair in Chinese hamster cells 4
 pleckstrin homology, Sec7 and coiled/coiled domains 4
 Consensus includes gp:AA47740 /FEA=EST /DB_XREF=gi:2161410 /DB_XREF=est:aa19105.s1 /CLONE=IMAGE.813633 /UG=Hs.99139 Homo sapiens cDNA: FLJ2294f:AA447740
 phosphodiesterase 4B, cAMP-specific (phosphodiesterase E4 duncce homolog, Drosophila)
 fatty acid binding protein 6, ileal (gastrotopin)
 translocase of outer mitochondrial membrane 70 homolog A (yeast)
 complement component 3a receptor 1
 gb:NM_005405.1 /DEF=Homo sapiens RNA binding motif protein, Y chromosome, family 2, member B (RBMV2B) mRNA. /FEA=mRNA /GEN=RBMV2B /PROD=RNA bindi
 hypothetical protein FLJ23119
 143 kDa E3 ubiquitin protein ligase (mouse)
 neural cell adhesion molecule 1
 KIAA0130 gene product
 Consensus includes gb:AW340588 /FEA=EST /DB_XREF=gi:6837214 /DB_XREF=est:hd05a05.x1 /CLONE=IMAGE.2906592 /UG=Hs.8736 Homo sapiens mRNA: cDNA
 heat shock protein (hsp110 family)
 hypothetical protein FLJ14393
 cer-d4 (mouse) homolog
 spectre-type POZ protein
 ras association (RalGDS/AF-6) domain containing protein JC265
 methionine adenosyltransferase II, beta
 Consensus includes gb:AB84887 /FEA=EST /DB_XREF=gi:5590031 /DB_XREF=est:wB5g03.x1 /CLONE=IMAGE.2431732 /UG=Hs.91379 ribosomal protein L26
 heterogeneous nuclear ribonucleoprotein U (scaffold attachment factor A)
 KIAA0275 gene product
 neuropilin 2
 PRO0159 protein
 tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)
 sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B
 putative leukocyte platelet-activating factor receptor
 Consensus includes gb:Y10255.1 /DEF=H.sapiens ACTH receptor mRNA 3UTR. /FEA=mRNA /DB_XREF=gi:2230860 /UG=Hs.274501 H.sapiens ACTH receptor mRNA 3'
 Y10259
 solute carrier family 21 (organic anion transporter), member 3
 cytotoxic T-lymphocyte-associated protein 4
 SH3-domain GRB2-like 3
 Consensus includes gb:AJ408433 /DEF=Homo sapiens partial IGKV gene for immunoglobulin kappa chain variable region, clone 38 /FEA=CDS /DB_XREF=gi:12655549 /AJ408433
 sine oculis homeobox homolog 2 (Drosophila)
 gap junction protein, beta 1, 32kD (connexin 32, Charcot-Marie-Tooth neuropathy, X-linked)
 DKFZP564I052 protein
 Consensus includes gb:AL137284.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434D1516 (from clone DKFZp434D1516). /FEA=mRNA /DB_XREF=gi:8807739 /UG=Hs.27
 tumor protein 63 kDa with strong homology to p53
 zinc-finger protein ZBRK1
 histone acetyltransferase
 aldehyde dehydrogenase 1 family, member A3
 retinoid X receptor, gamma
 karyopherin alpha 1 (importin alpha 5)
 protocadherin 22
 KIAA0652 gene product
 cathepsin H
 iduronidase, alpha-L-
 Consensus includes gb:AL520908 /FEA=EST /DB_XREF=gi:12784401 /DB_XREF=est:AL520908 /CLONE=CS0D8002YF01 (3 prime) /UG=Hs.155489 NS1-associated prc
 malonyl-CoA decarboxylase
 gap junction protein, alpha 5, 40kD (connexin 40)
 tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)

angiotensinogen 1
 cytochrome P450, subfamily 1 (dioxin-inducible), polypeptide 1 (glaucoma 3, primary infantile)
 WD repeat domain 3
 Z'-5'-oligoadenylate synthetase 3 (100 kD)
 annexin A13
 RAS guanyl releasing protein 2 (calcium and DAG-regulated)
 hypothetical protein FLJ21347
 mucin 1, transmembrane
 hypothetical protein FLJ12984
 carbonic anhydrase XII
 T-cell acute lymphocytic leukemia 1
 gb:AB058895.1 /DEF=Homo sapiens GHRHRps mRNA for growth hormone releasing hormone receptor, pituitary splice variant, complete cds. /FEA=mRNA /GEN=GHRH /AB058895
 similar to proline-rich protein 48
 KIAA1624 protein
 Consensus includes gb:AA865601 /FEA=EST /DB_XREF=gi:2957877 /DB_XREF=est:cg93603.s1 /CLONE=IMAGE:1455868 /UG=Hs.126281 Homo sapiens Chromosome
 a disintegrin and metalloproteinase domain 30
 casein, beta
 hypothetical protein FLJ20435
 solute carrier family 29 (nucleoside transporters), member 1
 soluble adenylyl cyclase
 dihydroxyimidine dehydrogenase
 lysosomal-associated protein transmembrane 4 alpha
 LIM domains containing 1
 SH3 protein interacting with Nck 90 kDa
 expressed in activated T/LAK lymphocytes
 hypothetical protein FLJ22029
 deleted in azoospermia-like
 gb:AJ225093.1 /DEF=Homo sapiens mRNA for single-chain antibody, complete cds (scFv2). /FEA=CDS /PROD=immunoglobulin /DB_XREF=gi:3090427 /UG=Hs.159386
 luciferiferase 3 (galactoside 3(4)-L-luciferiferase, Lewis blood group included)
 CD14 antigen
 superovulin
 adenovirus 5 E1A binding protein
 unc-13-like (C. elegans)
 osmosis responsive factor
 gb:NM_016528.1 /DEF=Homo sapiens hydroxyacid oxidase 3 (medium-chain) (HAO3), mRNA. /FEA=mRNA /GEN=HAO3 /PROD=hydroxyacid oxidase 3 /DB_XREF=gi:771 /NM_016528
 cold autoimmune syndrome 1
 ATPase, Class VI, type 11B
 Consensus includes gb:Y10259.1 /DEF=H sapiens ACTH receptor mRNA 3UTR. /FEA=mRNA /DB_XREF=gi:2230860 /UG=Hs.274501 H.sapiens ACTH receptor mRNA 3' UTR
 KIAA0844 protein
 KIAA1081 protein
 protein of P11, paired-like homeodomain transcription factor
 nuclear receptor coactivator 6
 hypothetical protein
 ectodysplasin A2 isoform receptor
 splicing factor 3a, subunit 2, 66kD
 DKFZP547E2110 protein
 early growth response 2 (Krox-20 homolog, Drosophila)
 RAB22A, member RAS oncogene family
 gonadotropin-releasing hormone 2
 keratin 2A (epidermal ichthyosis bullosa of Siemens)
 GREB1 protein
 Consensus includes gb:BE045982 /FEA=EST /DB_XREF=gi:8363035 /DB_XREF=est:hd91b04.x4 /CLONE=IMAGE:2916847 /UG=Hs.99093 Homo sapiens chromosome
 hypothetical protein FLJ21988
 chromosome 8 open reading frame 1
 ATP-binding cassette, sub-family C (CFTR/MRP), member 6
 protocadherin alpha 3
 potassium inwardly-rectifying channel, subfamily J, member 4
 DKFZP134D193 protein
 hypothetical protein
 hypothetical protein MGC16714
 potassium voltage-gated channel, Isk-related family, member 2
 inactivation escape 1
 Consensus includes gb:AK024135.1 /DEF=Homo sapiens cDNA FLJ14073 fs, clone HEMBB1001812. /FEA=mRNA /DB_XREF=gi:10436441 /UG=Hs.296751 Homo sapiens
 AK024135

proteoglycan 2, bone marrow (natural killer cell activator, eosinophil granule major basic protein) 1.17
 ATP-binding cassette, sub-family C (CFTR/MRP), member 9 1.17
 C-type lectin-like receptor-1 1.17
 a disintegrin and metalloproteinase domain 7 1.17
 zinc finger protein 281 1.17
 calmodulin 3 (phosphorylase kinase, delta) 1.17
 H4 histone family, member B 1.17
 Consensus includes gb:AK025271.1 /DEF=Homo sapiens cDNA: FLJ21618 fls, clone COL07487, /FEA=mRNA /DB_XREF=gi:10437753 /UG=Hs.306790 Homo sapiens cDNA: AK025271 1.17
 truncated repeat containing 11 (THR-associated protein, 230 kDa subunit) 1.17
 RAB2, member RAS oncogene family-like 1.17
 solute carrier family 16 (monocarboxylic acid transporters), member 5 1.17
 bromodomain adjacent to zinc finger domain, 1A 1.17
 small inducible cytokine subfamily A (Cys-Cys), member 24 1.17
 DKFZP586H2123 protein 1.17
 RBP2 interacting protein 8 1.17
 Consensus includes gb:AK025271.1 /DEF=Homo sapiens cDNA: FLJ21618 fls, clone COL07487, /FEA=mRNA /DB_XREF=gi:10437753 /UG=Hs.306790 Homo sapiens cDNA: AK025271 1.17
 hypothetical protein FLJ23451 1.17
 Consensus includes gb:AL049430.1 /DEF=Homo sapiens mRNA: cDNA DKFZb586H201 (from clone DKFZp586H201), /FEA=mRNA /DB_XREF=gi:4500210 /UG=Hs.3020 1.17
 dentatorubral-pallidolysian atrophy (atrophin-1) 1.17
 protocadherin gamma subfamily A, 11 1.17
 KIAA1609 protein 1.17
 cannabinoid receptor 2 (macrophage) 1.17
 capping protein (actin filament) muscle Z-line, alpha 2 1.17
 Consensus includes gb:AK024938.1 /DEF=Homo sapiens cDNA: FLJ21285 fls, clone COL01912, /FEA=mRNA /DB_XREF=gi:10437359 /UG=Hs.287655 Homo sapiens cDNA: AK024938 1.17
 hypothetical protein LOC37147 1.17
 retinoid X receptor, alpha 1.17
 SH2 domain protein 2A 1.17
 Consensus includes gb:BG34174 /FEA=EST /DB_XREF=gi:13340680 /DB_XREF=est:602506106F1 /CLONE=IMAGE:4603513 /UG=Hs.23437 Homo sapiens cDNA: FLJ1BG434174 1.17
 NK3 transcription factor homolog A (Drosophila) 1.17
 protein tyrosine phosphatase, receptor type, N polypeptide 2 1.17
 KIAA1237 protein 1.17
 kallikrein 1, renal/pancreas/salivary 1.17
 phosphodiesterase 6A, cGMP-specific, rod, alpha 1.17
 Consensus includes gb:Z39557 /FEA=EST /DB_XREF=gi:562749 /DB_XREF=est:HSC1DB062 /CLONE=c-1606 /UG=Hs.12533 Homo sapiens clone 23705 mRNA seque 239557 1.17
 transforming growth factor, alpha 1.17
 cyclin-dependent kinase-like 3 1.17
 Cluster Ind. AA522816/n40e12.a1 Homo sapiens cDNA, 3' end /clone=IMAGE:979342 /clone_end=3 /gb=AA522816 /gi=2263528 /ug=Hs.173789 /len=564 1.17
 partial transcript encompassing THC211630 gene 1.17
 similar to RIKEN cDNA 231004OG17 gene 1.17
 thionin oxidase 2 1.17
 cholinergic receptor, nicotinic, alpha polypeptide 4 1.17
 Consensus includes gb:AW974481 /FEA=EST /DB_XREF=gi:8165680 /DB_XREF=est:EST386585 /UG=Hs.293755 ESTs 1.17
 mesenchyme homeo box 1 1.17
 topomycin 4 1.17
 nuclear receptor subfamily 1, group H, member 3 1.17
 Consensus includes gb:AL535113 /FEA=EST /DB_XREF=gi:12798606 /DB_XREF=est:AL535113 /CLONE=CS00F008YC23 (3 prime) /UG=Hs.283006 phospholipase C, b 1.17
 KIAA1277 protein 1.17
 activator of CREM in testis 1.17
 Consensus includes gb:AK028106.1 /DEF=Homo sapiens cDNA: FLJ22453 fls, clone HRC09879, highly similar to AF059516 Homo sapiens telodiol-like 2 protein (TLL2) mR 1.17
 Consensus includes gb:AF035290.1 /DEF=Homo sapiens clone 23556 mRNA sequence, /FEA=mRNA /DB_XREF=gi:2661044 /UG=Hs.106300 Homo sapiens clone 23556 AF035290 1.17
 neuronal Shc adaptor homolog 1.17
 gamma-aminobutyric acid (GABA) A receptor, epsilon 1.17
 fasciculation and elongation protein zeta 2 (zyglin II) 1.17
 Mashapen/NIK-related kinase 1.17
 seizure related 6 homolog (mouse)-like 1.17
 low density lipoprotein receptor-related protein 6 1.17
 gb:NM_004250.1 /DEF=Homo sapiens RAB9, member RAS oncogene family, pseudogene 1 (RAB9P1), mRNA, /FEA=mRNA /GEN=RAB9P1 /PROD=RAB9, member RAS 1.17
 butyrophilin, subfamily 3, member A1 1.17
 Sho-20 related kinase 1.17
 interferon, alpha 6 1.17
 solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2 1.17
 Consensus includes gb:AA543076 /FEA=EST /DB_XREF=gi:2291556 /DB_XREF=est:n86102.s1 /CLONE=IMAGE:927771 /UG=Hs.82171 Homo sapiens (clone 115392) m 1.17
 Consensus includes gb:AA543076 /FEA=EST /DB_XREF=gi:2291556 /DB_XREF=est:n86102.s1 /CLONE=IMAGE:927771 /UG=Hs.82171 Homo sapiens (clone 115392) m 1.17

Consensus includes gb:AA716107 /FEA=EST /DB_XREF=est:zq63606.s1 /CLONE=IMAGE:398027 /UG=Hs.8963 Homo sapiens mRNA full length	1.17
topomycin 1 (alpha)	1.16
sperm associated antigen 1	1.16
collagen, type XV, alpha 1	1.17
phosphatidylinositol glycan, class B	1.17
histidine decarboxylase	1.17
Consensus includes gb:AW979289 /FEA=EST /DB_XREF=gij:8170577 /DB_XREF=est:EST3391399 /UG=Hs.293472 ESTs	1.16
gb:M73086.1 /DEF=Human androgen receptor mutant gene, mRNA, complete cds. /FEA=mRNA /GEN=AR /PROD=androgen receptor /DB_XREF=gij:178655 /FL=gb:M73086.1	1.16
hypoxia-inducible factor 1, alpha subunit inhibitor	1.16
secreted frizzled-related protein 1	1.16
hypothetical protein FLJ21820	1.16
putative non-coding transcript (DiGeorge critical region 5)	1.16
hypothetical protein FLJ20516	1.16
hypothetical protein FLJ23459	1.16
chromodomain helicase DNA binding protein 3	1.16
a distallegin and metalloproteinase domain 8	1.16
chromosome 11 open reading frame 16	1.16
myosin X	1.16
cysteine sulfinic acid decarboxylase-related protein 2	1.16
calponin like transmembrane domain protein	1.16
Consensus includes gb:BF224076 /FEA=EST /DB_XREF=gij:11131303 /DB_XREF=est:7q83a08.x1 /CLONE=IMAGE:3704942 /UG=Hs.307438 Human DNA sequence from BF224076	1.16
DnaJ (Hsp40) homolog, subfamily B, member 12	1.16
protein phosphatase 1A (formerly 2C), magnesium-dependent, alpha isoform	1.16
protocadherin beta 3	1.16
U5 snRNP-specific protein, 200 kDa (DEXH RNA helicase family)	1.16
Consensus includes gb:AA76303 /FEA=EST /DB_XREF=gij:2204514 /DB_XREF=est:zq28d09.s1 /CLONE=IMAGE:770705 /UG=Hs.306357 Homo sapiens isolate donor AA76303	1.16
hypothetical protein	1.16
gamma-aminobutyric acid (GABA) receptor, rho 2	1.16
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily f, member 1	1.16
dectonin	1.16
hypothetical protein DKFZp566A1524	1.16
KIAA0397 gene product	1.16
distal-less homeo box 5	1.16
nuclear protein 3 (apoptosis repressor with CARD domain)	1.16
Consensus includes gb:X07868 /DEF=Human DNA for insulin-like growth factor II (IGF-2); exon 7 and additional ORF /FEA=mRNA_1 /DB_XREF=gij:32998 /UG=Hs.25166r /X07868	1.16
transient receptor potential channel 1	1.16
KIAA0659 gene product	1.16
phosphatidylinositol 4-kinase, catalytic, beta polypeptide	1.16
hypothetical protein FLJ22559	1.16
gap junction protein, beta 3, 31kD (connexin 31)	1.16
olfactory receptor, family 1, subfamily F, member 2	1.16
hypothetical protein FLJ10759	1.16
Thy-1 cell surface antigen	1.16
scartin 6	1.16
Sec23 homolog A (S. cerevisiae)	1.16
hepatocellular carcinoma-associated gene TD28	1.16
HCGII-7 protein	1.16
deleted in colorectal carcinoma	1.16
interferon induced transmembrane protein 2 (1-8D)	1.16
vesicle-associated membrane protein 1 (synaptobrevin 1)	1.16
anti-Müllerian hormone	1.16
hypothetical protein FLJ13150	1.16
putative RhoG-interacting protein	1.16
MHC class I polypeptide-related sequence A	1.16
MAWD binding protein	1.16
mitogen-activated protein kinase kinase kinase 12	1.16
hypothetical protein EST00098	1.16
neurogenic differentiation 1	1.16
autoimmune regulator (autoimmune polyendocrinopathy candidiasis ectodermal dystrophy)	1.16
decay accelerating factor for complement (CD55, Crimer blood group system)	1.16
carbon-chloride cotransporter-interacting protein	1.16
KIAA01197 protein	1.16
epilepsy, progressive myoclonus type 2, Lafora disease (laforin)	1.16
KIAA02868	1.16

testis expressed sequence 15 NM_031271 1.16
zinc finger protein 42 (myeloid-specific retinoic acid- responsive) A173248 1.16
ribosomal protein L3-like NM_005061 1.16
cadherin 6, type 2, K-cadherin (fetal kidney) NM_004932 1.16
KIAA1548 protein NM_020909 1.16
scf-Coenzyme A oxidase 1, palmitoyl AB024518 1.16
DVS27-related protein NM_002721 1.16
Consensus includes gb:AK027217.1 /DEF=Homo sapiens cDNA: FLJ23564 fls, clone LNG10773. /FEA=mRNA /DB_XREF=gi:10440296 /UG=Hs.306913 Homo sapiens cC AK027217 AF225418 1.16
lipase protein NM_025033 1.16
hypothetical protein FLJ21277 H11725 1.16
KIAA1701 protein BC000103 1.16
NCK adaptor protein 2 AC005255 1.16
olfactory receptor, family 7, subfamily A, member 17 A1027678 1.16
KIAA0429 gene product NM_016284 1.16
KIAA1007 protein NM_003152 1.16
signal transducer and activator of transcription 5A NM_003152 1.16
Consensus includes gb:BE646315 /FEA=EST /DB_XREF=gi:9970626 /DB_XREF=est:764108.x1 /CLONE=IMAGE.3289191 /UG=Hs.292503 ESTs, Weakly similar to T471 BE646315 1.16
nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100) BC002844 1.16
Consensus includes gb:AK025064.1 /DEF=Homo sapiens cDNA: FLJ21411 fls, clone COL03986. /FEA=mRNA /DB_XREF=gi:10437503 /UG=Hs.306758 Homo sapiens cC AK025064 1.16
Consensus includes gb:AK025363.1 /DEF=Homo sapiens cDNA: FLJ21710 fls, clone COL10087. /FEA=mRNA /DB_XREF=gi:10437865 /UG=Hs.306807 Homo sapiens cC AK025363 A1346187 1.16
solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1 NM_001713 1.16
betaine-homocysteine methyltransferase NM_003318 1.16
TTK protein kinase AL503687 1.16
Consensus includes gb:AL583687 /FEA=EST /DB_XREF=gi:12952889 /DB_XREF=est:AL583687 /CLONE=CSDDJ008YL09 (5 prime) /UG=Hs.134852 ESTs NM_005711 1.16
EGF-like repeats and discoidin I-like domains 3 AL521391 1.16
ELL gene (11-19 lysine-rich leukemia gene) NM_020806 1.16
gephyrin M65217 1.16
heat shock transcription factor 2 NM_018952 1.16
homo box B6 AK022014 1.16
hypothetical protein FLJ11852 NM_001796 1.16
Consensus includes gb:AA1141166 /FEA=EST /DB_XREF=gi:1668052 /DB_XREF=est:zm75d3.s1 /CLONE=IMAGE.564004 /UG=Hs.23964 sin3-associated polypeptide, 18 AA1141166 1.16
KIAA0748 gene product NM_015644 1.16
DKFZP434B103 protein AL079292 1.16
Consensus includes gb:AK026682.1 /DEF=Homo sapiens cDNA: FLJ23029 fls, clone LNG01883. /FEA=mRNA /DB_XREF=gi:10435589 /UG=Hs.306864 Homo sapiens cC AK026682 1.16
hypothetical protein NM_018584 1.16
hypothetical protein PRO1489 A1692203 1.16
ring finger protein 3 NM_016642 1.16
spectrin, beta, non-erythrocytic 5 NM_012107 1.16
Consensus includes gb:AV761014 /FEA=EST /DB_XREF=gi:10918862 /DB_XREF=est:AV761014 /CLONE=MDSCCF09 /UG=Hs.291972 ESTs, Moderately similar to SC11 AV761014 1.16
retinoic acid induced 14 AF057557 1.16
regulator of Fas-induced apoptosis U44378 1.16
MAD, mothers against decapentaplegic homolog 4 (Drosophila) NM_000103 1.16
cytochrome P450, subfamily XIX (aromatization of androgens) NM_030931 1.16
epididymal secretory protein ESP13.2 AB004853 1.16
chromosome 21 open reading frame 18 U86495 1.16
leptin receptor NM_012107 1.16
bromodomain containing protein 75 kDa human homolog BF438056 1.16
cAMP responsive element binding protein-like 2 AF228413 1.16
nuclear receptor subfamily 5, group A, member 2 NM_024049 1.16
hypothetical protein MGC5566 NM_024049 1.16
Consensus includes gb:AK021578.1 /DEF=Homo sapiens cDNA FLJ11514 fls, clone HEMBA1002229. /FEA=mRNA /DB_XREF=gi:10432783 /UG=Hs.306802 Homo sapie AK021576 1.16
transforming growth factor beta-stimulated protein TSC-22 AK027071 1.16
matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase) NM_004994 1.16
Consensus includes gb:J04178.1 /DEF=Human abnormal beta-hexosaminidase alpha chain (HEXA) mRNA, partial cds. /FEA=mRNA /DB_XREF=gi:184001 /UG=Hs.1862 J04178 1.16
ring finger protein 5 BC004155 1.16
Rho guanine nucleotide exchange factor (GEF) 4 NM_015320 1.16
KIAA0286 gene product AA528252 1.16
KIAA0133 gene product NM_014777 1.16
NP0009 protein AF237813 1.16
hypothetical protein DKFZp41A023 AB037854 1.16
hypothetical protein FLJ13114 NM_024541 1.16
fibroblast growth factor receptor 3 (achondroplasia, thanatophoric dwarfism) M58051 1.16
endlease 3, (beta, muscle) NM_001976 1.16

gb:225427.1 /DEF=H.sapiens protein-serine/threonine kinase gene, complete CDS, /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405732 /FL=gb:225. Z25427
 claudin 1 NM_021101 1.16
 galactosylceramidase (Krabbe disease) D25284 1.16
 GA binding protein transcription factor, beta subunit 1 (53kD) NM_005254 1.16
 CSE1 chromosome segregation 1-like (yeast) AF053841 1.16
 hypothetical protein FLJ13725 NM_024519 1.16
 KIAA0892 protein AW157731 1.16
 hypothetical protein FLJ20216 NM_017715 1.16
 myelin transcription factor 1 M86980 1.16
 protein tyrosine phosphatase, receptor type, A BF740139 1.16
 nuclear receptor binding protein NM_013392 1.16
 EphA3 AF213460 1.16
 peptidylglycine alpha-amidating monooxygenase NM_000919 1.16
 carboxypeptidase D BE349147 1.16
 signal recognition particle 54kD NM_003136 1.16
 imprinted in Prader-Willi syndrome AI672541 1.16
 RAN binding protein 8 NM_006390 1.16
 musashi homolog 1 (Drosophila) NM_002442 1.16
 tumor necrosis factor receptor superfamily, member 4 AJ277151 1.16
 dentin, light polypeptide (Lca) NM_001833 1.16
 solute carrier family 12 (potassium/chloride transporters), member 4 U55054 1.16
 Consensus includes gb:U55987 /FEA=EST /DB_XREF=est:HSU55987 /CLONE=47507 /UG=Hs.323773 ESTs. Highly similar to NPC1_HUMAN NIE U55987 1.16
 ATP-binding cassette, sub-family A (ABC1), member 7 NM_019112 1.16
 transducin (beta)-like 2 AK026529 1.16
 2',5'-oligoadenylate synthetase 1 (40-46 kD) NM_018816 1.16
 Consensus includes gb:AL046983.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564D042. /FEA=mRNA /DB_XREF=gi:4884234 /UG=Hs.2401 AL046983 1.16
 hypothetical protein FLJ20051 NM_018067 1.16
 Cluster ind. A1248730q440b01.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:1871401 /clone_end=3 /gb=A1246730 /gi=3842127 /ug=Hs.126901 /len=770 A1246730 1.16
 hypothetical protein A-21108.1 BE963444 1.16
 myosin, light polypeptide 5, regulatory NM_022477 1.16
 spilloprotein B48 receptor NM_018690 1.16
 monocyte to macrophage differentiation-associated NM_012328 1.16
 reelin NM_005045 1.16
 ELK4, ETS-domain protein (SRF accessory protein 1) NM_021795 1.16
 cholecystokinin B receptor BC000740 1.16
 B7 protein NM_008892 1.16
 lectin, galactoside-binding, soluble, 8 (galectin 8) AF342815 1.16
 EGF-TM7-180philin-related protein NM_022159 1.16
 N-sulfoglucosamine sulfohydrolase (sulfamidase) NM_000199 1.16
 coagulation factor II (thrombin) receptor-like 2 NM_004101 1.16
 nuclear receptor subfamily 1, group H, member 2 NM_007121 1.16
 bryotin-like NM_004053 1.16
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4 AA772023 1.16
 KIAA1031 protein NM_025013 1.16
 interleukin 22 NM_020525 1.16
 hypothetical protein FLJ21868 NM_022769 1.16
 lung cancer candidate NM_002725 1.16
 glutathione S-transferase M4 NM_000850 1.16
 olfactory receptor, family 7, subfamily C, member 2 NM_012377 1.16
 hypothetical protein FLJ20359 NM_017781 1.16
 putative DNA binding protein BG033764 1.16
 FYN binding protein (FYB-120/130) AF198052 1.16
 piccolo (presynaptic cytomatrix protein) BC001304 1.16
 RAB7, member RAS oncogene family-like 1 BG338251 1.16
 nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1 U08015 1.16
 islet cell autoantigen 1 (89kD) NM_022307 1.16
 macrophage scavenger receptor 1 NM_002445 1.16
 chromosome 19 open reading frame 3 NM_005716 1.16
 fibronectin 1 X02761 1.16
 NY-REN-24 antigen BC004282 1.16
 KIAA0040 gene product NM_014656 1.16
 solute carrier family 22 (organic anion transporter), member 7 AF210455 1.16
 latent transforming growth factor beta binding protein 2 NM_000428 1.16

LIM homeobox protein 2 U11701 1.16
 synovial sarcoma translocation, chromosome 18 AF257501 1.16
 Consensus includes gb:BE409994 /FEA=EST /DB_XREF=gi:9346444 /DB_XREF=est:601299404F1 /CLONE=IMAGE:3629592 /UG=Hs.326464 Homo sapiens cDNA: FLJ: BE409994 1.16
 immunoglobulin superfamily, member 3 AB007935 1.16
 G protein-coupled receptor 75 NM_006794 1.16
 3-hydroxytryptophan 3,4-dioxygenase NM_012205 1.16
 Consensus includes gb:H94842 /FEA=EST /DB_XREF=gi:1102475 /DB_XREF=est:y57a03.31 /CLONE=IMAGE:230188 /UG=Hs.17882 ESTs H94842 1.16
 phospholipase A2, group IVA (cytosolic, calcium-dependent) M88874 1.16
 D13642 1.16
 A1625402 1.16
 NM_002256 1.16
 NM_003281 1.16
 NM_021645 1.16
 NM_017512 1.16
 rTS beta protein AL049381 1.16
 Consensus includes gb:AL049381.1 /DEF=Homo sapiens mRNA: cDNA DKFZp586J2118 (from clone DKFZp586J2118). /FEA=mRNA /DB_XREF=gi:4500168 /UG=Hs.218 AL049381 1.16
 VENT-like homeobox 2 NM_018065 1.16
 hypothetical protein FLJ10346 M24779 1.16
 p1m-1 oncogene NM_014652 1.16
 importin 13 NM_016134 1.16
 plasma glutamate carboxypeptidase NM_018019 1.16
 Consensus includes gb:BF111846 /FEA=EST /DB_XREF=gi:10941536 /DB_XREF=est:7036001.x1 /CLONE=IMAGE:3523608 /UG=Hs.106334 Homo sapiens clone 23636 BF111846 1.16
 hypothetical protein FLJ10193 AK026980 1.16
 Consensus includes gb:AK026980.1 /DEF=Homo sapiens cDNA: FLJ23327 fls, clone HEP12630, highly similar to HSNF37 Homo sapiens ZNF37A mRNA for zinc finger f AK026980 1.16
 hypothetical protein FLJ23191 NM_024574 1.16
 HTGN29 protein NM_020199 1.16
 polymerase (DNA directed), theta NM_014125 1.16
 neuropeptide FF 1; RFamide-related peptide receptor NM_022146 1.16
 mitogen-activated protein kinase kinase 4 NM_006724 1.16
 ovarian cancer overexpressed 1 NM_015945 1.16
 ATP-binding cassette, sub-family A (ABC1), member 2 AF327657 1.16
 ethandamine kinase NM_018638 1.16
 ligand of neuronal nitric oxide synthase with carboxyl-terminal PDZ domain AF037070 1.16
 amiloride-sensitive cation channel 1, neuronal (degenerin) NM_001094 1.16
 Consensus includes gb:H53689 /FEA=EST /DB_XREF=gi:993836 /DB_XREF=est:y136g07.r1 /CLONE=IMAGE:236124 /UG=Hs.306239 Homo sapiens clone ASPBLL54 in H53689 1.16
 peptidylarginine deiminase type 1 AK023188 1.16
 protein phosphatase 1, regulatory (inhibitor) subunit 13B NM_013358 1.16
 hypothetical protein FLJ14281 NM_024920 1.16
 interleukin 5 receptor, alpha M86651 1.16
 KIAA0853 protein, B7-like protein AF199028 1.16
 soluble carrier family 8 (neurotransmitter transporter, betaine/GABA), member 12 U27699 1.16
 leucine zipper, putative tumor suppressor 1 N21184 1.16
 interleukin 13 receptor, alpha 1 NM_001560 1.16
 hypothetical protein FLJ23071 NM_025192 1.16
 v-jun sarcoma virus 17 oncogene homolog (avian) NM_002228 1.16
 adenylate cyclase activating polypeptide 1 (pituitary) NM_001117 1.16
 chemokine (C-C motif) receptor 7 NM_001838 1.16
 T cell receptor alpha locus AE000659 1.16
 Consensus includes gb:AL162040.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434N199 (from clone DKFZp434N199). /FEA=mRNA /DB_XREF=gi:7328079 /UG=Hs.3064 AL162040 1.16
 BCL2-like 11 (apoptosis facilitator) NM_008538 1.16
 AD024 protein AF225418 1.16
 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2 NM_025190 1.16
 stanniocalcin 1 U46768 1.16
 cleavage and polyadenylation specific factor 6, 68kD subunit NM_007007 1.16
 lymphocyte antigen 84 homolog, activating NK-receptor; NK-p46, (mouse) AJ008121 1.16
 GL-Kruppel family member GL13 (Grieg cephalopolysyndactyly syndrome) NM_000168 1.16
 KIAA0014 gene product NM_014685 1.16
 Consensus includes gb:BC004327.1 /DEF=Homo sapiens, clone IMAGE:3629896. mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE:3629896) /DB_XI BC004327 1.16
 KIAA1841 protein NM_025190 1.16
 zinc finger protein 144 (Met-18) NM_007144 1.16
 Consensus includes gb:AK022094.1 /DEF=Homo sapiens cDNA FLJ12032 fls, clone HEMBB1001880. /FEA=mRNA /DB_XREF=gi:10433415 /UG=Hs.151194 Homo sapiens AK022094 1.16
 phospholipase A2, group IIE NM_014589 1.16
 Consensus includes gb:AL109707.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 295344. /FEA=mRNA /DB_XREF=gi:5889824 /UG=Hs.943271 AL109707 1.16
 Consensus includes gb:L41657 /DEF=Homo sapiens synthetic myelin basic protein 21.5 kDa isoform gene, complete cds /FEA=cds /DB_XREF=gi:1162921 /UG=Hs.2480 L41657 1.16

v-erb-b2 erythroblastic leukemia viral oncogene homolog 2, neurofiblioblastoma derived oncogene homolog (avian)		
guanylate cyclase activator 1A (telina)	AF177761	1.16
HMB-Inducible	L36861	1.16
protein tyrosine phosphatase type IVA, member 1	AW665096	1.16
mitogen-activated protein kinase 14	AL578310	1.16
phorodin-like protein MDS019	L35253	1.16
rab3 GTPase-activating protein, non-catalytic subunit (150kD)	BF508948	1.16
leukocyte tyrosine kinase	BF240652	1.16
growth arrest-specific 7	X52213	1.16
hypothetical protein FLJ20156	BE439987	1.16
hypothetical protein FKSG28	NM_017691	1.16
KIAA0382 protein	NM_030929	1.16
laminin, beta 2 (laminin S)	A1348094	1.16
thyroid hormone receptor, alpha (erythroblastic leukemia viral (v-erb-a) oncogene homolog, avian)	X79683	1.16
hypothetical protein FLJ20244	NM_003250	1.16
actin-like 6	AL390133	1.16
flightless I homolog (Drosophila)	NM_016188	1.16
tumor protein D52	U80184	1.16
hypothetical protein FLJ13159	AA524023	1.16
KIAA0082 protein	NM_021940	1.16
CD163 antigen	D43949	1.16
LPS-induced TNF-alpha factor	Z22970	1.16
v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog	AB034747	1.16
S100 calcium binding protein A2	W80678	1.16
hypothetical protein DKF2p564F013	NM_030927	1.16
retinoic acid receptor, gamma	NM_005978	1.16
CGI-01 protein	AL050389	1.16
KIAA0013 gene product	M57707	1.16
glutaredoxin 2	AK001172	1.16
crystallin, zeta (quinone reductase)	NM_014783	1.16
chromosome 9 open reading frame 9	NM_016066	1.16
Consensus includes gb:AK000244.1 /DEF=Homo sapiens cDNA FLJ20237 fs, clone COLF5879, highly similar to AB023230 Homo sapiens mRNA for KIAA1013 protein, fl	NM_001889	1.16
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)	NM_018956	1.15
leukocyte receptor cluster (LRC) member 5	M61213	1.15
LCA1-like lysophospholipase	NM_002010	1.15
Consensus includes gb:AI446234 /FEA=EST /DB_XREF=gi:4293782 /DB_XREF=est:j25g10.x1 /CLONE=IMAGE:2142594 /UG=Hs.156072 Human pre-TNK cell associated	NM_024075	1.15
diolipinase, iodotyrosine, type II	AL110209	1.15
spondin 1, (I-spondin) extracellular matrix protein	AB041844	1.15
hypothetical protein DKF2p34L0850	AB016305	1.15
potassium inwardly-rectifying channel, subfamily J, member 6	NM_017558	1.15
zinc finger protein 73 (Cos12)	N39314	1.15
ubiquitin specific protease 1	NM_012480	1.15
glycoprotein 2 (zymogen granule membrane)	AW489935	1.15
T cell receptor gamma constant 2	NM_016295	1.15
CD86 antigen (CD28 antigen ligand 2, B7-2 antigen)	AL580863	1.15
tumor necrosis factor, alpha-induced protein 1 (endothelial)	NM_017559	1.15
KIAA1451 protein	NM_006889	1.15
matrix metalloproteinase 11 (stromelysin 3)	BC001643	1.15
hippocalcin-like protein 4	AL048923	1.15
glycophorin B (includes Ss blood group)	AL136591	1.15
forkhead box J1	U05255	1.15
hypothetical protein FLJ21749	NM_001454	1.15
Gardner-Rasheed feline sarcoma viral (v-igr) oncogene homolog	NM_025124	1.15
retinaldehyde binding protein 1	NM_065248	1.15
KIAA0275 gene product	NM_000326	1.15
deleted in esophageal cancer 1	A1952009	1.15
RAR (RAS like GTPASE)	NM_017418	1.15
estrogen receptor 1	BE965869	1.15
Nijmegen breakage syndrome 1 (nibrin)	NM_000125	1.15
CD2 antigen (p50), sheep red blood cell receptor	A1766269	1.15
	NM_001767	1.15

Consensus includes gb:AW675473 /FEA=EST /DB_XREF=gi:7540699 /DB_XREF=est:bs49g05.x1 /CLONE=IMAGE:2899928 /UG=Hs.101916 Homo sapiens mRNA; cDN / AW675473
CDC14 cell division cycle 14 homolog A (S. cerevisiae) 1.15
daudin 3 AF084102 1.15
hypothetical protein FLJ22378 NM_025078 1.15
chloride intracellular channel 3 BE791251 1.15
ATP-binding cassette, sub-family B (MOT/TAP), member 1 NM_004669 1.15
apolipoprotein E AF016535 1.15
hypothetical protein FLJ21032 N33009 1.15
Carbonic anhydrase-related protein 10 NM_024906 1.15
lymphocyte antigen 117 NM_020178 1.15
transient receptor potential channel 5 AF031138 1.15
brain and nasopharyngeal carcinoma susceptibility protein NM_012471 1.15
Consensus includes gb:AU148274 /FEA=EST /DB_XREF=gi:11009795 /DB_XREF=est:AU148274 /CLONE=MAMMA1003089 /UG=Hs.301201 Homo sapiens cDNA FLJ14 /AU148274 1.15
putative transmembrane protein AF191492 1.15
polycystic kidney disease 2-like 2 NM_007011 1.15
sulfotransferase family, cytosolic, 1C, member 1 NM_014386 1.15
lymphocyte antigen 95 (activating NK-receptor; NK-p44) AF186255 1.15
cycochrome P450, subfamily IIA, polypeptide 7 NM_004828 1.15
NY-REN-24 antigen AF315325 1.15
zinc finger protein A013329 1.15
putative receptor (family A group 5) NM_014415 1.15
aquaporin 4 NM_005767 1.15
CD38 antigen (collagen type I receptor, thrombospondin receptor) U83622 1.15
small membrane protein 1 NM_016382 1.15
hypothetical protein FLJ14360 NM_000072 1.15
nuclear RNA export factor 2 NM_014313 1.15
MAX protein N38751 1.15
kallikrein 11 NM_017809 1.15
gb:BC006370.1 /DEF=Homo sapiens. Similar to inversin, clone MGC:12888, mRNA, complete cds. /FEA=mrna /PROD=Similar to inversin /DB_XREF=gi:13623522 /FL-g
doublecortin; lissencephaly, X-linked (doublecortin) NM_006853 1.15
polymerase (RNA) II (DNA directed) polypeptide A (220kD) BC006370 1.15
androgen receptor-associated coregulator 267 AF040254 1.15
neurotrophin M21610 1.15
Consensus includes gb:AU160041 /FEA=EST /DB_XREF=gi:1021562 /DB_XREF=est:AU160041 /CLONE=Y79AA1000989 /UG=Hs.120858 Homo sapiens cDNA FLJ139 /AU160041 1.15
heparan sulfatase (glucosaminase) 3-O-sulfotransferase 1 AW117456 1.15
HLA-B associated transcript 2 NM_003460 1.15
Consensus includes gb:AA72320 /FEA=EST /DB_XREF=gi:4334410 /DB_XREF=est:407c02.x1 /CLONE=IMAGE:2148482 /UG=Hs.48504 ESTs. Moderately similar to ALU /AA72320 1.15
zinc finger protein 135 (clone pTZ-17) NM_003436 1.15
kinesin family member 5C BF059313 1.15
Consensus includes gb:AL109682.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 35394. /FEA=mrna /DB_XREF=gi:5685804 /UG=Hs.23777 H /AL109682 1.15
zona pellucida glycoprotein 2 (sperm receptor) NM_003460 1.15
phosphorylase kinase, beta NM_000293 1.15
leukocyte receptor cluster (LRC) member 4 BC003164 1.15
chromogranin B (secretogranin 1) NM_001819 1.15
Dystonia musculorum of mouse, human homolog of AL049215 1.15
histone H2A.F/Z variant A4534860 1.15
prostate androgen-regulated transcript 1 AT770098 1.15
serine dehydratase NM_006843 1.15
collagen, type IX, alpha 2 AF733465 1.15
interleukin 13 receptor, alpha 1 U81360 1.15
phosphodiesterase 4D, cAMP-specific (phosphodiesterase E3 dunce homolog, Drosophila) U50157 1.15
mitogen-activated protein kinase kinase 6 NM_004672 1.15
interleukin 8 AF043337 1.15
Consensus includes gb:AV763520 /FEA=EST /DB_XREF=gi:10921368 /DB_XREF=est:AV763520 /CLONE=MDSCH03 /UG=Hs.282113 ESTs
cadherin 5, type 2 (T1-cadherin) NM_016278 1.15
nuclear factor of kappa light polypeptide gene enhancer in B-cells 2 (p49/p100) NM_002502 1.15
KIAA1133 protein AL037338 1.15
KIAA1045 protein AB028968 1.15
sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F AL138552 1.15
DKF ZP566C0424 protein AL582808 1.15
putative integral membrane transporter AW149681 1.15
Consensus includes gb:AL137457.1 /DEF=Homo sapiens mRNA; cDNA DKF ZP566G231.1 (from clone DKF ZP566G231). /FEA=mrna /DB_XREF=gi:6808035 /UG=Hs.2723 AL137457

F-box only protein 5
Cluster Inci: AB06793:wf15405.x1 Homo sapiens cDNA, 3' end (clone=IMAGE:2350665 (done_end=3 /gb=AB06793 /gi=5393359 /ug=Hs.239450 /len=555
neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)
hypothetical protein FLJ20375
interleukin 15 receptor, alpha
hypothetical protein FLJ11274
spinocerebellar ataxia 1 (olivopontocerebellar ataxia 1, autosomal dominant, ataxia 1)
KIAA1055 protein
hypothetical protein FLJ23506
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9
insulin receptor
hypothetical protein MQC5580
SRY (sex determining region Y-box 5
daudin 11 (oligodendrocyte transmembrane protein)
hypothetical protein FLJ12387 similar to kinesin light chain
hypothetical protein FLJ20309
homeodomain interacting protein kinase 3
hypothetical protein PRO2949
aldehyde dehydrogenase 1 family, member B1
Consensus includes gb:AB041269.1 /DEF=Homo sapiens mRNA for keratin 19, partial cds, isolate:K19-141 /FEA=mRNA /GEN=keratin 19 /PROD=keratin 19 /DB_XREF=gi:3368883 /UG=Hs.2638 /AL390149
ankyrin 2, neuronal
hypothetical protein FLJ13511
KIAA1037 protein
H1 histone family, member 4
protein phosphatase 2, regulatory subunit B (B56), beta isoform
CCAAT-box-binding transcription factor
UDP-N-acetylglucosamine-2-epimerase/N-acetylmannosamine kinase
adenylyate kinase 5
prion protein (PrP^{Sc}) (Creutzfeldt-Jakob disease, Gerstmann-Sträussler-Scheinker syndrome, fatal familial insomnia)
steroid-5-desaturase (fungal ERG3, delta-5-desaturase)-like
small inducible cytokine subfamily A (Cys-Cys), member 21
leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 5
signal transducer and activator of transcription 1, 91KD
hypothetical protein FLJ11160
TATA box binding protein (TBP)-associated factor, RNA polymerase II, S, 150KD
adipose most abundant gene transcript 1
KIAA0728 gene product
phosphatidylethanolamine N-methyltransferase
stromal cell-derived factor 1
triple helix repeat containing 21
spondin 2, extracellular matrix protein
sulfite oxidase
HSPC054 protein
KIAA0391 gene product
KIAA1028 protein
putative Ly-6 superfamily member
hypothetical protein PRO0689
solute carrier family 21 (organic anion transporter), member 11
Consensus includes gb:AL390149.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547B026 (from clone DKFZp547B026) /FEA=mRNA /DB_XREF=gi:3368883 /UG=Hs.2638 /AL390149
a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 6
glutaminase
STAT induced STAT inhibitor-4
calcium/calmodulin-dependent protein kinase IG
egl-like module containing, much-like, hormone receptor-like sequence 2
synaptosomal complex protein 1
proteoglycan gamma subfamily B, 5
regulator of G-protein signalling 4
Consensus includes gb:AF070591.1 /DEF=Homo sapiens clone 24649 mRNA sequence /FEA=mRNA /DB_XREF=gi:3387968 /UG=Hs.322473 Homo sapiens clone 24649 AF070591
cathepsin B
cAMP responsive element binding protein-like 2
RE1-silencing transcription factor
Consensus includes gb:U63542.1 /DEF=Human putative FAP protein mRNA, partial cds. /FEA=mRNA /PROD=FAP protein mRNA /DB_XREF=gi:1890646 /UG=Hs.166101 Human U63542
KIAA1097 protein

triple functional domain (PTPRF interacting)
 kalikrein 13
 N-acetylglucosaminidase, alpha-
 CULL8 protein
 cholinergic receptor, nicotinic, alpha polypeptide 5
 protein tyrosine phosphatase, receptor type, R
 hypothetical protein FLJ11726
 Consensus includes gb:AK026484.1 /DEF=Homo sapiens cDNA, FLJ22831 fls, clone KIAA1461, /FEA=mRNA /DB_XREF=gi:10439356 /UG=Hs.321666 Homo sapiens cDI AK026484
 salivary proline-rich protein
 melanoma inhibitory activity
 transforming, acidic coiled-coil containing protein 1
 peptidyl-prolyl isomerase G (cyclophilin G)
 class I cytokine receptor
 stromal antigen 2
 complement component (3d/Epstein Barr virus) receptor 2
 Consensus includes gb:AU147528 /FEA=EST /DB_XREF=gi:11009049 /DB_XREF=est:AU147528 /CLONE=MAMMA1000940 /UG=Hs.301811 Homo sapiens cDNA FLJ14
 Consensus includes gb:AK024138.1 /DEF=Homo sapiens cDNA FLJ14076 fls, clone HEMBB1001825, /FEA=mRNA /DB_XREF=gi:10439445 /UG=Hs.306667 Homo sapie
 tropoin 1, cardiac
 KIAA0218 gene product
 CGI-78 protein
 protein C receptor, endothelial (EPCR)
 Kruppel-like factor 7 (ubiquitous)
 ubiquitously transcribed tetratricopeptide repeat gene, X chromosome
 Consensus includes gb:AK024381.1 /DEF=Homo sapiens cDNA FLJ14319 fls, clone PLACE3000406, /FEA=mRNA /DB_XREF=gi:10436753 /UG=Hs.287638 Homo sapie
 Consensus includes gb:BE672181 /FEA=EST /DB_XREF=gi:10032712 /DB_XREF=est:7b51c08.x1 /CLONE=IMAGE:3231758 /UG=Hs.155489 NS1-associated protein 1 /F
 interferon, gamma-inducible protein 16
 prostaglandin E receptor 3 (subtype EP3)
 SGC32445 protein
 core-binding factor, runt domain, alpha subunit 2; translocated to 1: cyclin D-related
 potassium voltage-gated channel, delayed-rectifier, subfamily S, member 1
 cut-like 1, CCAAT displacement protein (Drosophila)
 hypothetical protein FLJ20055
 hypothetical protein FLJ22638
 G protein-coupled receptor, family C, group 5, member B
 AUP-ribosylarginine hydrolase
 Consensus includes gb:AF070620.1 /DEF=Homo sapiens clone 24694 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3283886 /UG=Hs.158530 Homo sapiens clone 24694
 thyroid hormone receptor-associated protein, 150 kDa subunit
 glycine dehydrogenase (decarboxylating; glycine decarboxylase, glycine cleavage system protein P)
 isovaleryl Coenzyme A dehydrogenase
 Consensus includes gb:AU144792 /FEA=EST /DB_XREF=gi:11006313 /DB_XREF=est:AU144792 /CLONE=HEMBA1002973 /UG=Hs.319540 Homo sapiens cDNA FLJ10
 caspase recruitment domain family, member 15
 mannosyl (alpha-1,3)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
 G protein-coupled receptor 19
 septin 8
 hypothetical protein FLJ10083
 Consensus includes gb:AU146655 /FEA=EST /DB_XREF=gi:11008176 /DB_XREF=est:AU146655 /CLONE=HEMBA1001133 /UG=Hs.288178 Homo sapiens cDNA FLJ11
 Consensus includes gb:BF062299 /FEA=EST /DB_XREF=gi:10821197 /DB_XREF=est:7N76c03.x1 /CLONE=IMAGE:3481325 /UG=Hs.323079 Homo sapiens mRNA; cDN
 delta-tubulin
 KIAA0657 protein
 Fc fragment of IgG, low affinity IIb, receptor for (CD32)
 NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 6 (14kD, B14)
 microtubule-associated protein 4
 ankyrin 1, erythrocytic
 cocaine- and amphetamine-regulated transcript
 purinergic receptor P2X, ligand-gated ion channel, 2
 obaferrin
 SRY (sex determining region Y)-box 22
 glucosaminyl (N-acetyl) transferase 3, mucin type
 wingless-type MMTV integration site family member 2
 hypothetical protein FLJ23312
 KIAA1078 protein
 solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 2
 hypothetical protein FLJ2724

Consensus includes gb:BF508835 /FEA=EST /DB_XREF=gi:11592133 /DB_XREF=est:U1-H-B14-acc-d-03-0-JUI.s1 /CLONE=IMAGE:3085853 /UG=Hs.6217 Homo sapiens (BF508835) 1.15
twisted gastrulation 1.15
deleted in azoospermia 1.15
CD3G antigen, gamma polypeptide (T1T3 complex) 1.15
Consensus includes gb:X61072.1 /DEF=Human mRNA for T cell receptor, clone IGRA17. /FEA=mRNA /PROD=T cell receptor /DB_XREF=gi:33511 /UG=Hs.272500 Huma X61072 1.15
Consensus includes gb:X61072.1 /DEF=Human mRNA for T cell receptor, clone IGRA17. /FEA=mRNA /PROD=T cell receptor /DB_XREF=gi:33511 /UG=Hs.272500 Huma X61072 1.15
COP22 for nondatrhin coat protein zeta-COP 1.15
Conserved gene telomeric to alpha globin cluster 1.15
dopa decarboxylase (aromatic L-amino acid decarboxylase) 1.15
cancer/testis antigen 2 1.15
CCR4-NOT transcription complex, subunit 4 1.15
KIAA1085 protein 1.15
thioredoxin reductase 1 1.15
hypothetical protein MGC10784 1.15
phorbol-like protein MDS019 1.15
Consensus includes gb:BE504197 /FEA=EST /DB_XREF=gi:9706605 /DB_XREF=est:h6611.x1 /CLONE=IMAGE:3180332 /UG=Hs.129061 Human Chromosome 16 BA1 BE504197 1.15
kallikrein 3, (prostate specific antigen) 1.15
5-hydroxytryptamine (serotonin) receptor 1E 1.15
hypothetical protein FLJ10842 1.15
phospholipase D1, phosphatidylcholine-specific 1.15
sprouty homolog 2 (Drosophila) 1.15
brain-specific angiogenesis inhibitor 1 1.15
alpha-2-macroglobulin 1.15
protein tyrosine phosphatase, receptor type, I polypeptide (PTPRF), interacting protein (liprin), alpha 2 1.15
paired box gene 8 1.15
caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2) 1.15
sarcosine dehydrogenase 1.15
EphB2 1.15
period homolog 3 (Drosophila) 1.15
Consensus includes gb:S81916.1 /DEF=phosphoglycerate kinase (alternatively spliced) human, phosphoglycerate kinase deficient patient with episodes of muscle 1.15
major histocompatibility complex, class I-like sequence 1.15
inositol 1,4,5-trisphosphate receptor, type 1 1.15
hypothetical protein FLJ20199 1.15
hypothetical protein MGC10715 1.15
phosphatidylinositol-4-phosphate 5-kinase, type 1, gamma 1.15
Consensus includes gb:AL049301.1 /DEF=Homo sapiens mRNA; cDNA DKFZ564P073 (from clone DKFZ564P073). /FEA=mRNA /DB_XREF=gi:4500067 /UG=Hs.2785 AL049301 1.15
neuropeptide Y receptor Y5 1.15
ras homolog gene family, member 8 1.15
hypothetical protein FLJ20551 1.15
HCGIV-6 protein 1.15
statmin-like 3 1.15
hypothetical protein FLJ22173 1.15
Consensus includes gb:AL356954 /DEF=Human DNA sequence from clone RP11-520F24 on chromosome 13 Contains ESTs, STSs and GSSs. Contains an HNRPA1 (het 1.15
RNA binding motif protein 9 1.15
hypothetical protein FLJ11184 1.15
homolog of mouse quaking OKI (KH domain RNA binding protein) 1.15
glycoprotein M6A 1.15
Consensus includes gb:AK026679.1 /DEF=Homo sapiens cDNA: FLJ23026 fls, clone LNG01738. /FEA=mRNA /DB_XREF=gi:10439585 /UG=Hs.308663 Homo sapiens cD AK026679 1.15
hypothetical protein FLJ10850 1.15
KIAA1446 protein 1.15
SET binding factor 1 1.15
retinoic acid receptor responder (tazarotene induced) 3 1.15
ribosomal protein L37a 1.14
dopamine receptor D2 1.14
hypothetical protein 1.14
Consensus includes gb:BG485135 /FEA=EST /DB_XREF=gi:13417414 /DB_XREF=est:602503756F1 /CLONE=IMAGE:4617445 /UG=Hs.325722 Immunoglobulin kappa v 1.14
cullin 5 1.14
plestin C1 1.14
syndecin 1 1.14
gb:NM_016374.2 /DEF=Homo sapiens RBP1-like protein (BCAA). mRNA. /FEA=mRNA /GEN=BCAA /PROD=RBP1-like protein /DB_XREF=gi:13518237 /FL=gb:NM_016374 1.14
angiotensin-like 3 1.14
Consensus includes gb:AK024849.1 /DEF=Homo sapiens cDNA: FLJ21196 fls, clone COL00193. /FEA=mRNA /DB_XREF=gi:10437257 /UG=Hs.287850 Homo sapiens cD AK024849 1.14
hypothetical protein MGC3279 similar to collectins 1.14
kuronidase, alpha-L- 1.14

collagen, type VI, alpha 1
angiotensin 1 converting enzyme (peptidyl-dipeptidase A) 2
Consensus includes gb:134409.1 /DEF=Homo Sapiens (clone 83B3E13) chromosome 4p16.3 DNA fragment. /FEA=mRNA /DB_XREF=gi:600517 /UG=Hs.94799 Homo Sapiens
butyrophilin, subfamily 1, member A1
Consensus includes gb:AL133388 /DEF=Human DNA sequence from clone RP1-181C24 on chromosome 6p11.1-12.2. Contains the 3' end of the BMP5 gene for bone morphogenetic protein kinase domain 19 (meltrin beta)
a disintegrin and metalloproteinase component 2
progesterone receptor membrane component 2
thioredoxin reductase 2
carboxyl ester lipase (bile salt-stimulated lipase)
neuridin 1
gb:L14455.1 /DEF=Human Ig rearranged mu-chain gene V-N-D-N-J-reion, complete cds. /FEA=mRNA /DB_XREF=gi:398496 /FL=gb:L14455.1
TEA domain family member 4
hypothetical protein FLJ22386
prip28, US snRNP 100 kd protein
glutamate decarboxylase 1 (brain, 67kD)
contactin associated protein-like 2
trypsin gamma 1
prostaglandin E receptor 3 (subtype EP3)
RAB31, member RAS oncogene family
dachshund homolog (Drosophila)
hypothetical protein B
MAP/microtubule affinity-regulating kinase like 1
YME1-like 1 (S. cerevisiae)
poly(A) polymerase beta (testis specific)
zinc finger protein 207
metallothionein-like 5, testis-specific (testin)
arachidonate lipooxygenase 3
transcription termination factor, RNA polymerase I
Consensus includes gb:AL109716.2 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROMIMAGE 208948. /FEA=mRNA /DB_XREF=gi:9187597 /UG=Hs.200341
dual specificity phosphatase 6
Consensus includes gb:AF052148.1 /DEF=Homo sapiens clone 24507 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360456 /UG=Hs.169744 Homo sapiens clone 24507
hypothetical protein FLJ11726
hypothetical protein FLJ10488
Consensus includes gb:BF970253 /FEA=EST /DB_XREF=gi:12337468 /DB_XREF=est:602273505F1 /CLONE=IMAGE:4361569 /UG=Hs.173684 Homo sapiens mRNA; cDNA DKFZ547E184 (from clone DKFZ547E184). /FEA=mRNA /DB_XREF=gi:9368866 /UG=Hs.8077
zinc finger protein 200
hypothetical protein FLJ11560
plexin B2
gb:NM_020887.1 /DEF=Homo sapiens hypothetical protein from clone 1659351 (LOC57397), mRNA. /FEA=mRNA /GEN=LOC57397 /PROD=hypothetical protein from clone NM_020887
solute carrier family 22 (organic cation transporter), member 1
myosin, heavy polypeptide 11, smooth muscle
hypothetical protein FLJ10490
splicing factor 3a, subunit 2, 86kD
carbonic anhydrase XI
Rho guanine nucleotide exchange factor (GEF) 4
Consensus includes gb:AL049544 /DEF=Human DNA sequence from clone 24088 on chromosome 6p11.2-q12. Contains the 3' part of a gene for a novel protein similar to melanoma antigen, family A, 3
Consensus includes gb:AL390170.1 /DEF=Homo sapiens mRNA; cDNA DKFZ547E184 (from clone DKFZ547E184). /FEA=mRNA /DB_XREF=gi:9368866 /UG=Hs.8077
melanoma antigen, family A, 3
Consensus includes gb:BF110411 /FEA=EST /DB_XREF=gi:10940101 /DB_XREF=est:7n5202.x1 /CLONE=IMAGE:3568274 /UG=Hs.16959 ESTs
SPFR protein kinase 2
Cluster Inc. A1417917:ig55d04.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2112679 /clone_end=3 /gb=A1417917 /gi=4261421 /ug=Hs.20733 /len=662
regulator of G-protein signalling 5
neuronal pentraxin 1
potassium channel, subfamily K, member 2 (TREK-1)
collagen, type V, alpha 1
dynactin 4
ATP synthase, H+ transporting, mitochondrial F1 complex, epsilon subunit
KIAA1504 protein
sialic acid binding Ig-like lectin 8
HRI/HFB2206 protein
ATP synthase, H+ transporting, mitochondrial F1 complex, gamma polypeptide 1
popay protein 3
eukaryotic translation termination factor 1

AA292373 1.14
AK026461 1.14
NM_001732 1.14
NM_002756 1.14
NM_023038 1.14
BE879873 1.14
BF513089 1.14
NM_001807 1.14
NM_014932 1.14
L14455 1.14
NM_003213 1.14
NM_024589 1.14
NM_004818 1.14
NM_000817 1.14
AC005378 1.14
NM_012467 1.14
X83858 1.14
NM_006868 1.14
AW772082 1.14
NM_014262 1.14
AL120554 1.14
NM_014263 1.14
NM_020144 1.14
NM_003457 1.14
NM_004923 1.14
AW003512 1.14
AB32304 1.14
AL109716 1.14
BC003143 1.14
NM_024971 1.14
NM_018110 1.14
BF970253 1.14
AU148447 1.14
AL583909 1.14
AL022328 1.14
NM_020887 1.14
NM_003057 1.14
AB89739 1.14
NM_018111 1.14
BC004434 1.14
AB018195 1.14
AB042199 1.14
AL049544 1.14
AL390170 1.14
BC000340 1.14
BF110411 1.14
NM_003138 1.14
A1417917 1.14
AF169570 1.14
NM_002522 1.14
AF004711 1.14
AB93428 1.14
BC004191 1.14
NM_006886 1.14
NB4622 1.14
AF287892 1.14
W19873 1.14
BG232034 1.14
AB928544 1.14
NM_004730 1.14

TONDU
 glutamate rich WD repeat protein GRWD
 HYAZ2 protein
 tumor necrosis factor alpha-inducible cellular protein containing leucine zipper domains: Huntingtin Interacting protein L: transcription factor IIIA-interacting protein
 Consensus includes gb:AK000993.1 /DEF=Homo sapiens cDNA FLJ10131 fls, clone HEMBA1003041 /FEA=mRNA /DB_XREF=gi:7021996 /UG=Hs.274128 Homo sapiens dual specificity phosphatase 10
 protein tyrosine phosphatase, non-receptor type substrate 1
 Consensus includes gb:AA757368 /FEA=EST /DB_XREF=gi:2805231 /DB_XREF=est:ah98a10.s1 /CLONE=IMAGE:1327098 /UG=Hs.7057 Homo sapiens Crl-du-chat regl A4757368
 coagulation factor C (Limulus polyphemus homolog), coctilin
 Consensus includes gb:AA343027 /FEA=EST /DB_XREF=gi:1995263 /DB_XREF=est:EST46865 /UG=Hs.193606 Homo sapiens PAC clone RP5-109301.7 from 7q11.23-q AA343027
 Consensus includes gb:AA343027 /FEA=EST /DB_XREF=gi:1995263 /DB_XREF=est:EST46865 /UG=Hs.193606 Homo sapiens PAC clone RP5-109301.7 from 7q11.23-q AA343027
 protocadherin 22
 Consensus includes gb:AU148161 /FEA=EST /DB_XREF=gi:1009682 /DB_XREF=est:AU148161 /CLONE=MAMMA1002758 /UG=Hs.296720 Homo sapiens cDNA FLJ12 AU148161
 Interleukin 1 receptor, type I
 KIAA0328 protein
 NS1-binding protein
 KIAA0971 protein
 folate receptor 1 (adult)
 Consensus includes gb:AL139318 /DEF=Human DNA sequence from clone RP11-124B17 on chromosome 13. Contains the DCT gene for dopachrome tautomerase (dopa AL139318
 hypothetical protein
 ubiquitin ligase factor E4A (UFD2 homolog, yeast)
 human homolog of Drosophila Scribble
 phosphatidic acid phosphatase type 2B
 hypothetical protein FLJ14251
 Consensus includes gb:U09902.1 /DEF=Human clone 23612 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1913880 /UG=Hs.82141 Human clone 23612 mRNA sequence U09902
 UOP-Gal-beta-D-GlcNAc beta 1,4-galactosyltransferase, polypeptide 1
 KIAA1065 protein
 leucine-rich repeat protein, neuronal 1
 a disintegrin and metalloproteinase domain 5
 Consensus includes gb:AA338945 /FEA=EST /DB_XREF=gi:1991203 /DB_XREF=est:EST43994 /UG=Hs.318825 Homo sapiens, clone IMAGE:3616574, mRNA, partial cd A4338945
 angiotensin 1
 hypothetical protein
 DNFP366F1524 protein
 Consensus includes gb:AK025152.1 /DEF=Homo sapiens cDNA: FLJ21499 fls, clone COL05634. /FEA=mRNA /DB_XREF=gi:10437612 /UG=Hs.306773 Homo sapiens cDNA AK025152
 glutamate receptor, ionotropic, N-methyl D-aspartate 2B
 nucleosome assembly protein 1-like 3
 hypothetical protein FLJ13544
 Consensus includes gb:AK025363.1 /DEF=Homo sapiens cDNA: FLJ21710 fls, clone COL10087. /FEA=mRNA /DB_XREF=gi:10437865 /UG=Hs.306807 Homo sapiens cDNA AK025363
 mesh homeo box homolog, 1 (Drosophila)
 hypothetical protein FLJ10116
 Consensus includes gb:AA176809 /FEA=EST /DB_XREF=gi:2820147 /DB_XREF=est:nz82h01.s1 /CLONE=IMAGE:1302001 /UG=Hs.293205 ESTs, Weakly similar to BC3 AA768909
 splicing factor, arginine/serine-rich 5
 hypothetical protein FLJ10496
 mannan-binding lectin serine protease 2
 HGC8.1.1 protein
 hair/enhancer-of-split related with YRPW motif 2
 N-acetylglucosamine amidohydrolase (acid ceramidase)
 core1 UDP-galactose 4-epimerase
 Consensus includes gb:BF987657 /FEA=EST /DB_XREF=gi:12334872 /DB_XREF=est:3597895 /DB_XREF=est:34402.x1 /CLONE=IMAGE:1711491 /UG=Hs.89463 potassium large conductance c A1729381
 5-hydroxytryptamine (serotonin) receptor 1F
 hypothetical protein FLJ11275
 matrix metalloproteinase 9 (gelatinase B, 92kD gelatinase, 92kD type IV collagenase)
 centromere protein A (17kD)
 Consensus includes gb:AW974915 /FEA=EST /DB_XREF=gi:8166118 /DB_XREF=est:EST3787020 /UG=Hs.116550 ESTs
 Consensus includes gb:AI129381 /FEA=EST /DB_XREF=gi:3597895 /DB_XREF=est:34402.x1 /CLONE=IMAGE:1711491 /UG=Hs.89463 potassium large conductance c A1729381
 KIAA0237 gene product
 Cluster Ind. N31716:yy15c12.s1 Homo sapiens cDNA, 3 end /clone=IMAGE:271318 /clone_end=3 /gb=N31716 /gi=1152115 /ug=Hs.31754 /len=624
 SH3-domain binding protein 5 (BTK-associated)
 hyperpolarization activated cyclic nucleotide-gated potassium channel 4
 hypothetical protein FLJ22557
 clone 1900 unknown protein
 cysteine dioxygenase, type I
 hypothetical protein FLJ20300

aryl hydrocarbon receptor nuclear translocator-like
KIAA0790 protein
Consensus includes gb:AU155621 /FEA=EST /DB_XREF=gi:11017142 /DB_XREF=est:AU155621 /CLONE=PLACE1000444 /UG=Hs.306418 Homo sapiens cDNA FLJ108 AU155621
hypotheical protein FLJ10120
caudal type homeo box transcription factor 4
hypotheical protein FLJ10726
hypotheical protein
Consensus includes gb:AL137428.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761N1323 (from clone DKFZp761N1323) /FEA=mRNA /DB_XREF=gi:5807885 /UG=Hs.30 AL137428
hypotheical gene FLJ13072
unc-51-like kinase 1 (C. elegans)
interleukin 11
eukaryotic translation initiation factor 1A
lymphoid enhancer-binding factor 1
fibronectin 1
Consensus includes gb:AL049997.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564L102 (from clone DKFZp564L102) /FEA=mRNA /DB_XREF=gi:4884250 /UG=Hs.3063C AL049997
tumor necrosis factor receptor superfamily, member 6b, decoy
brain cell membrane protein 1
tubby super-family protein
actin related protein 2/3 complex, subunit 4 (20 kD)
similar to ubiquitin binding protein
sperm associated antigen 11
cardioembryonic antigen-related cell adhesion molecule 3
Consensus includes gb:A1732905 /FEA=EST /DB_XREF=gi:5054018 /DB_XREF=est:ot72a12.x5 /CLONE=IMAGE:1435870 /UG=Hs.184507 Homo sapiens Chromosome 1 A1732905
dica, large (Drosophila) homolog-associated protein 2
S164 protein
defensin, alpha 5, Peneth cell-specific
glutamate decarboxylase 2 (pancreatic islets and brain, 65kD)
CDC14 cell division cycle 14 homolog A (S. cerevisiae)
transmembrane 4 superfamily member 1
translocase of inner mitochondrial membrane 17 homolog B (yeast)
Charot-Leyden crystal protein
splicing factor 3b, subunit 3, 130kD
G protein-coupled receptor 51
gb:AF116705.1 /DEF=Homo sapiens PRO2489 mRNA, complete cds. /FEA=mRNA /PROD=PRO2489 /DB_XREF=gi:7959908 /FL=gb:AF116705.1
ubiquitin-conjugating enzyme E2 variant 1
hypotheical protein MGC2821
Consensus includes gb:U79304.1 /DEF=Human clone 23909 mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1710291 /UG=Hs.281043 Human clone 239 U79304
X-ray repair complementing defective repair in Chinese hamster cells 2
matrix metalloproteinase 11 (stromelysin 3)
prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)
DiGeorge syndrome critical region gene DGS1; likely ortholog of mouse expressed sequence 2 embryonic lethal
golgi SNAP receptor complex member 2
Consensus includes gb:AK021690.1 /DEF=Homo sapiens cDNA FLJ11628 fis, clone HEMBA1004238. /FEA=mRNA /DB_XREF=gi:10432926 /UG=Hs.306610 Homo sapiens AK021690
killer cell lectin-like receptor subfamily F, member 1
zinc transporter
DKFZP434D146 protein
Rhesus blood group, CcEe antigens
Mets1, myeloid ecotropic viral integration site 1 homolog (mouse)
hypotheical protein FLJ22389
hypotheical protein FLJ20433
zinc finger protein 136 (clone pHZ-20)
Consensus includes gb:AL390143.1 /DEF=Homo sapiens mRNA; cDNA DKFZp762C115 (from clone DKFZp762C115) /FEA=mRNA /DB_XREF=gi:9368878 /UG=Hs.2843 AL390143
BCM-like membrane protein precursor
schaefer-scute complex-like 1 (Drosophila)
Consensus includes gb:U00956.1 /DEF=Human clone KDB5.1 (CACAC)(GTG)n repeat-containing mRNA. /FEA=mRNA /DB_XREF=gi:405058 /UG=Hs.104112 Human clone U00956
synuclein, beta
putative selenocysteine lyase
hypotheical protein FLJ20211
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1
lecithin-cholesterol acyltransferase
promyelocytic leukemia
hypotheical protein FLJ10671
Consensus includes gb:AL109730.1 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIIMAGE 68600. /FEA=mRNA /DB_XREF=gi:5689835 /UG=Hs.306331 AL109730

v-fos FBJ murine osteosarcoma viral oncogene homolog
hypothetical protein ES100098
caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)
Consensus includes gb:AL137673.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434H0872 (from clone DKFZp434H0872), /FEA=mRNA /DB_XREF=gi:680784.1 /UG=Hs.30 AL137673
interferon-induced protein with tetrapeptide repeats 4
hypothetical protein FLJ14297
DKFZP56850519 protein
lysophospholipase II
calbindin 2, (29kD, calretinin)
absent in melanoma 2
protein tyrosine phosphatase, receptor type, N polypeptide 2
Consensus includes gb:A1732802 /FEA=EST /DB_XREF=gi:5053915 /DB_XREF=est:ab23e08.x5 /CLONE=IMAGE:841670 /UG=Hs.292679 ESTs
hypothetical protein FLJ14154
collagen, type XIV, alpha 1 (undulin)
HSPC072 protein
synaptophysin-like protein
cytochrome b-5
hypothetical protein FLJ12154
EphB6
neurogenic differentiation 4
KIAA1609 protein
H2B histone family, member Q
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 1
DKFZP5682420 protein
POU domain, class 3, transcription factor 1
leucyl-RNA synthetase, mitochondrial
KIAA0674 protein
eyes absent homolog 3 (Drosophila)
Consensus includes gb:AL049883.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564D042 (from clone DKFZp564D042), /FEA=mRNA /DB_XREF=gi:4884234 /UG=Hs.2401 AL049883
transducin (beta)-like 1
hypothetical protein FLJ20449
poly(A) polymerase gamma
truncal repeat containing 1
COX11 homolog, cytochrome c oxidase assembly protein (yeast)
pvi-1 (murine) oncogene homolog, MYC activator
baculoviral IAP repeat-containing 1
KIAA0346 protein
Consensus includes gb:AL354915 /DEF=Human DNA sequence from clone RP11-392A19 on chromosome 13. Contains ESTs, STSs and GSSs. Contains a novel pseudogene AL354915
hair/enhancer-oligo-split related with YRPW motif-like
KIAA0669 gene product
hypothetical protein similar to preferentially expressed antigen of melanoma
cyclin H
FK506 binding protein 1A (12kD)
Consensus includes gb:AF203728.1 /DEF=Homo sapiens bone morphogenetic protein BMPY (BMPY) mRNA, partial cds. /FEA=mRNA /GEN=BMPY /PROD=bone morpho AF203728
KIAA1058 protein
ATP-dependent RNA helicase
ANKHZN protein
hypothetical protein MGC2592
ATP-binding cassette, sub-family C (CFTR/MRP), member 2
Consensus includes gb:U82303.1 /DEF=Homo sapiens unknown protein mRNA, partial cds. /FEA=mRNA /PROD=unknown /DB_XREF=gi:1938329 /UG=Hs.123080 Homo U82303
matrix metalloproteinase 24 (membrane-inserted)
hypothetical protein DKFZp434E0321
osylator 7 alpha-hydroxylase
N-myc (and STAT) interactor
Consensus includes gb:AF189251.1 /DEF=Homo sapiens cytomegalovirus partial fusion receptor mRNA, partial cds. /FEA=mRNA /PROD=cytomegalovirus partial fusion r AF189251
G4 protein
gamma-aminobutyric acid (GABA) receptor, theta
sphingosine kinase 1
transition protein 2 (during histone to protamine replacement)
hypothetical protein FLJ11939
T cell receptor alpha locus
KIAA0092 gene product
solute carrier family 7, (cationic amino acid transporter, y+ system) member 11
NM_014331

Consensus includes gb:A8935123 /FEA=EST /DB_XREF=gi:5673993 /DB_XREF=est:wp13n09.x1 /CLONE=IMAGE:2464769 /UG=Hs.57548 ESTs 1.14

CD89 antigen (p80, early T-cell activation antigen) 1.14

hypothetical protein FLJ12888 1.14

cyclin D2 1.14

ZW10 interactor 1.14

POU domain, class 5, transcription factor 1 1.14

Tax1 (human T-cell leukemia virus type 1) binding protein 1 1.14

tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor) 1.14

gamma tubulin ring complex protein (76p gene) 1.14

aminopeptidase puromycin sensitive 1.14

diuronate 2-sulfatase (Hunter syndrome) 1.14

Consensus includes gb:AK024231.1 /DEF=Homo sapiens cDNA FLJ14169 fts, clone NT2RP2002056 /FEA=mRNA /DB_XREF=gi:10435558 /UG=Hs.306670 Homo sapiens 1.14

ring finger protein 1 1.14

karyopherin alpha 3 (importin alpha 4) 1.14

REV1-like (yeast) 1.14

plasmolipin 1.14

KIAA0614 protein 1.14

myosin, heavy polypeptide 4, skeletal muscle 1.14

interleukin 2 receptor, alpha 1.14

Consensus includes gb:A879383 /FEA=EST /DB_XREF=gi:5553432 /DB_XREF=est:au50b05.x1 /CLONE=IMAGE:2518161 /UG=Hs.190885 ESTs, Moderately similar to A A1879383 1.14

promyelocytic leukemia 1.14

proteoglycan 1, secretory granule 1.14

Consensus includes gb:AK022686.1 /DEF=Homo sapiens cDNA FLJ12824 fts, clone NT2RM44001754 /FEA=mRNA /DB_XREF=gi:10434225 /UG=Hs.308639 Homo sapiens 1.13

chromosome 1 open reading frame 27 1.13

kidney- and liver-specific gene 1.13

PRO1598 protein 1.13

translation factor su11 homolog 1.13

collagen, type IV, alpha 6 1.13

hyaluronan binding protein 2 1.13

angiotensin-like 2 1.13

chemokine (C-C motif) receptor 9 1.13

KIAA0076 gene product 1.13

putative translation initiation factor 1.13

seven in absentia homolog 1 (Drosophila) 1.13

hypothetical protein 1.13

Kv channel interacting protein 2 1.13

hypothetical protein FLJ20331 1.13

trophinin 1.13

RAS p21 protein activator (GTPase activating protein) 3 (ins(1,3,4,5)p4-binding protein) 1.13

erlaine homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila) 1.13

apoptosis-associated tyrosine kinase 1.13

add cluster protein 33 1.13

amyloid beta (A4) precursor protein (protease nexin-II, Alzheimer disease) 1.13

dual specificity phosphatase 1 1.13

hypothetical protein FLJ11795 1.13

KIAA0354 gene product 1.13

Consensus includes gb:BG231551 /FEA=EST /DB_XREF=gi:12726658 /DB_XREF=est:na129h05.x1 /CLONE=IMAGE:4142457 /UG=Hs.74861 activated RNA polymerase I BG231551 1.13

KIAA0763 gene product 1.13

double C2-like domains, beta 1.13

luteinizing hormone/choriogonadotropin receptor 1.13

bromodomain-containing 1 1.13

pyrimidinergic receptor P2Y, G-protein coupled, 6 1.13

keratin 13 1.13

EHM2 gene 1.13

hypothetical protein FLJ21511 1.13

adenosine deaminase 1.13

oncomodulin 1.13

hypothetical protein FLJ23233 1.13

protein tyrosine phosphatase, non-receptor type 21 1.13

ATP-binding cassette, sub-family C (CFTR/MRP), member 5 1.13

beta-amyloid binding protein precursor 1.13

hypothetical protein FLJ20753 1.13

myoneurium 1.13

A8935123 1.14

L07555 1.14

NM_024945 1.14

NM_001759 1.14

NM_007057 1.14

NM_002701 1.14

U25801 1.14

AF021834 1.14

BF592982 1.14

AW340850 1.14

BC006170 1.14

AK024231 1.14

NM_002931 1.14

AL120704 1.14

NM_016316 1.14

AL020869 1.14

BC006270 1.14

NM_017533 1.14

NM_000417 1.14

A1879383 1.14

BC000080 1.14

J03223 1.14

AK022686 1.13

NM_017847 1.13

AB013094 1.13

NM_031270 1.13

NM_005875 1.13

AL031177 1.13

NM_004132 1.13

AF007150 1.13

AF145439 1.13

D38548 1.13

AL523310 1.13

NM_003031 1.13

NM_015371 1.13

NM_014591 1.13

NM_017788 1.13

AF349720 1.13

NM_007368 1.13

N25546 1.13

NM_004920 1.13

NM_016630 1.13

X06989 1.13

NM_004417 1.13

NM_024669 1.13

NM_014872 1.13

NM_014869 1.13

NM_003585 1.13

NM_000233 1.13

NM_014577 1.13

NM_004154 1.13

NM_002274 1.13

NM_019114 1.13

NM_025087 1.13

Z97053 1.13

NM_006188 1.13

NM_024691 1.13

X79510 1.13

AF146074 1.13

AA012917 1.13

NM_017950 1.13

NM_018657 1.13

Consensus includes gb:AJ278883 /DEF=Homo sapiens alpha-NAC gene for nascent polypeptide-associated complex component /FEA=CDS /DB_XREF=gi:9968289 /UG= AJ278883
 bone morphogenetic protein receptor, type IA NM_004329
 kunronate 2-sulfatase (Hunter syndrome) AF179281
 cyclin-dependent kinase 2 M68520
 protein kinase, lysine deficient 1 A1788512
 KIAA0602 protein AL583340
 astrotactin AL031290
 dihydrofolamide branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex, maple syrup urine disease) J03208
 UDP-Gal4betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 4 AB026730
 solute carrier family 12 (sodium/chloride transporters), member 3 NM_000339
 SRY (sex determining region Y)-box 10 A1367319
 placenta-specific 1 NM_021796
 toll-like receptor 5 AF051151
 A1868626
 HUS1 checkpoint homolog (S. pombe) NM_017450
 BAI1-associated protein 2 NM_024014
 Consensus includes gb:N92708 /FEA=EST /DB_XREF=gi:1265017 /DB_XREF=est:zb44c04.s1 /CLONE=IMAGE:306438 /UG=Hs.32374 ESTs, Weakly similar to KIAA152t N92708
 homeo box A6 NM_016170
 NCX protein AA527502
 heterogeneus nuclear ribonucleoprotein A3 NM_001870
 carbonic dehydratase A3 (mast cell) AF043977
 chloride channel, calcium activated, family member 2
 Consensus includes gb:AV759552 /FEA=EST /DB_XREF=gi:10917400 /DB_XREF=est:AV759552 /CLONE=MDSBCH06 /UG=Hs.226138 Homo sapiens mRNA; cDNA DK1 AV759552
 diacylglycerol kinase, zeta (104kD) Z81010
 platelet-derived growth factor beta polypeptide (simian sarcoma viral (v-sis) oncogene homolog) AF191654
 nudix (nucleoside diphosphate linked moiety X)-type motif 4 DB1792
 PBX1/notch 1 homeobox 2
 Consensus includes gb:BE669609 /FEA=EST /DB_XREF=gi:10030150 /DB_XREF=est:7a15h11.x1 /CLONE=IMAGE:3282597 /UG=Hs.251411 Homo sapiens mRNA; cDN BE669609
 Consensus includes gb:AK025239.1 /DEF=Homo sapiens cDNA: FLJ21586 fls, clone COL06920. /FEA=mRNA /DB_XREF=gi:10437709 /UG=Hs.321142 Homo sapiens cDNA AK025239
 hypothetical protein FLJ11457 NM_024753
 lectin, mannose-binding, 1 NM_005570
 peptidylarginine deiminase type III NM_016233
 alpha-1,4-N-acetylglucosaminyltransferase NM_016161
 myosin binding protein C, fast type NM_004533
 protocadherin gamma subfamily A, 3 AF152509
 protein kinase Njmu-R1 NM_022344
 Consensus includes gb:BE669609 /FEA=EST /DB_XREF=gi:10030150 /DB_XREF=est:7a15h11.x1 /CLONE=IMAGE:3282597 /UG=Hs.251411 Homo sapiens mRNA; cDN BE669609
 Consensus includes gb:AK024615.1 /DEF=Homo sapiens cDNA: FLJ20962 fls, clone ADH00804. /FEA=mRNA /DB_XREF=gi:10436931 /UG=Hs.306698 Homo sapiens cDNA AK024615
 Consensus includes gb:U79277.1 /DEF=Human done 23548 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710245 /UG=Hs.71846 Human done 23548 mRNA sequence U79277
 small nuclear RNA activating complex, polypeptide 4, 190kD AK023513
 desmulin AK026420
 hypothetical protein FLJ12903 NM_022753
 I-box and leucine-rich repeat protein 2 AF176518
 phosphatidylinositol-4-phosphate 5-kinase, type II, beta NM_003559
 Consensus includes gb:Ala28075 /FEA=EST /DB_XREF=gi:5446746 /DB_XREF=est:wk28c04.x1 /CLONE=IMAGE:2413538 /UG=Hs.297681 serine (or cysteine) proteinase A1826075
 Consensus includes gb:A075181 /FEA=EST /DB_XREF=gi:3401772 /DB_XREF=est:oy96b04.x1 /CLONE=IMAGE:1673647 /UG=Hs.304000 ESTs, Highly similar to inosito A075181
 ATPase, Cu++ transporting, beta polypeptide (Wilson disease) NM_000053
 paired box gene 6 (aniridia, keratitis) NM_000280
 S100 calcium binding protein A14 (calgizarin) NM_021039
 interleukin 17B NM_014443
 peptidylprolyl isomerase (cyclophilin)-like 2 BC000022
 homeo box B1 S49765
 D component of complement (cd1psin) NM_001928
 amyloid beta (A4) precursor-like protein 1 U48437
 oxidative 3 alpha hydroxysteroid dehydrogenase; retinol dehydrogenase; 3-hydroxysteroid epimerase U89281
 prostaglandin I2 (prostaglandin) synthase D38145
 Consensus includes gb:AU148808 /FEA=EST /DB_XREF=gi:11010328 /DB_XREF=est:AU148808 /CLONE=NT2RM4000979 /UG=Hs.96927 Homo sapiens cDNA FLJ125t; AU148808
 KIAA0847 protein AW294686
 Consensus includes gb:A1820796 /FEA=EST /DB_XREF=gi:5439875 /DB_XREF=est:an35h08.x5 /CLONE=IMAGE:1700703 /UG=Hs.310806 ESTs, Moderately similar to A A1820796
 thymidylate synthetase BG281679
 deleted in azoospermia AF271088
 Consensus includes gb:AL024493 /DEF=Human DNA sequence from clone 452M16 on chromosome Xq21.1-21.33 Contains capping protein alpha subunit isoform 1 pseud AL024493
 semaphorin sem2 NM_020163
 Cluster incl. A4225490.nc25b01.r1 Homo sapiens cDNA /clone=IMAGE:1009129 /gb=AA225490 /gi=1846817 /ug=Hs.80115 /len=539 A4225490

golgii-like protein	NM_018652	1.13
apurinic/apyrimidinic endonuclease(APEX nuclease)-like 2 protein	NM_014481	1.13
hypothetical protein PRO2730	AL137854	1.13
major histocompatibility complex, class II, DQ beta 1	M17955	1.13
DKFZP568N0319 protein	NM_015437	1.13
regulator of G-protein signaling 13	AF030107	1.13
hypothetical protein FLJ11598	NM_024966	1.13
KIAA0451 gene product	NM_014828	1.13
Consensus includes gb:U80773.1 /DEF=Human EST clone 42944 mariner transposon Hamar1 sequence. /FEA=mRNA /DB_XREF=gi:121580 Human ES U80773	NM_000194	1.13
hypoxanthine phosphoribosyltransferase 1 (Lesch-Nyhan syndrome)	BE983245	1.13
I-box and WD-40 domain protein 1B	BC003617	1.13
RAB6A, member RAS oncogene family	NM_006657	1.13
formin/nucleation factor 1 (Lesc-Nyhan syndrome)	NM_014442	1.13
sialic acid binding Ig-like lectin 8	U92226	1.13
myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))	NM_006322	1.13
spindle pole body protein	NM_021622	1.13
pleckstrin homology domain-containing, family A (phosphoinositide binding specific) member 1	U42349	1.13
Pituitary prostate cancer tumor suppressor	U82532	1.13
Rho GDP dissociation inhibitor (GDI) gamma	BC003683	1.13
cytochrome-like nuclear factor n-pac	BE971373	1.13
cytochrome P450, subfamily 11C (mephenytoin 4-hydroxylase), polypeptide 9	AE596111	1.13
KIAA0653 protein, 87-like protein	NM_003305	1.13
transient receptor potential channel 3	NM_021942	1.13
Consensus includes gb:BF574664 /FEA=EST /DB_XREF=gi:11648376 /DB_XREF=est:602132139F1 /CLONE=IMAGE:4271792 /UG=Hs.274531 Homo sapiens mRNA; cDNA	BF574664	1.13
vascular endothelial growth factor C	U58111	1.13
Consensus includes gb:AK025325.1 /DEF=Homo sapiens cDNA: FLJ21672 fa, clone COL09025. /FEA=mRNA /DB_XREF=gi:10437818 /UG=Hs.306801 Homo sapiens cDNA	NM_004313	1.13
arrestin, beta 2	NM_004909	1.13
taxol resistance associated gene 3	NM_058336	1.13
Consensus includes gb:AA583336 /FEA=EST /DB_XREF=gi:2367945 /DB_XREF=est:tm45108.s1 /CLONE=IMAGE:1086855 /UG=Hs.27371 Homo sapiens mRNA; cDNA	AA583336	1.13
hypothetical protein FLJ12716	NM_004504	1.13
HIV-1 Rev binding protein	AL041139	1.13
jerky homolog (mouse)	NM_002347	1.13
lymphocyte antigen 6 complex, locus H	BC002715	1.13
peroxisome proliferative activated receptor, delta	NM_002782	1.13
pregnancy specific beta-1-glycoprotein 6	AW305097	1.13
phosphatidylinositol glycan, class C, pseudogene 1	AA694291	1.13
SON DNA binding protein	AB013384	1.13
huntingtin interacting protein-1-related	AF220532	1.13
nuclear receptor subfamily 2, group E, member 1	NM_016616	1.13
NM23-H8	AF140507	1.13
calcium/calmodulin-dependent protein kinase 2, beta	AL136868	1.13
hypothetical protein DKFZP434P144	NM_004694	1.13
Consensus includes gb:AV704017 /FEA=EST /DB_XREF=gi:10721338 /DB_XREF=est:AV704017 /CLONE=ADBAOG04 /UG=Hs.15833 ESTs, Moderately similar to ALU7.	AV704017	1.13
solute carrier family 16 (monocarboxylic acid transporters), member 6	AY009401	1.13
wingless-type MMTV integration site family, member 6	NM_007250	1.13
Kruppel-like factor 6	NM_024764	1.13
hypothetical protein FLJ14298	AB044343	1.13
UDP-glucuronic acid/UDP-N-acetylglucosamine dual transporter	NM_006898	1.13
homeo box A7	BC003167	1.13
phosphatidylinositol 4-kinase type II	NM_002061	1.13
Consensus includes gb:U63332.1 /DEF=Human super cysteine rich protein mRNA, partial cds. /FEA=mRNA /PROD=super cysteine rich protein /DB_XREF=gi:1480862 UC	U63332	1.13
glutamate-cysteine ligase, modifier subunit	BF11821	1.13
DKFZP564A122 protein	NM_014156	1.13
DKFZP564O0463 protein	NM_003951	1.13
solute carrier family 25 (mitochondrial carrier, brain), member 14	NM_003813	1.13
cartilage intermediate layer protein, nucleotide pyrophosphatase	NM_000558	1.13
hemoglobin, alpha 1	BC005325	1.13
synovial sarcoma, X breakpoint 4	NM_001825	1.13
creatine kinase, mitochondrial 2 (sarcomeric)	NM_003807	1.13
tumor necrosis factor (ligand) superfamily, member 14	BF433902	1.13
tumor necrosis factor receptor superfamily, member 11b (osteoprotegerin)	AF082537	1.13
culin 3	NM_022569	1.13
N-deacetylaseN-sulfotransferase 4		

NAG-7 protein
 keratocan
 hypothetical protein PRO2133
 mitogen-activated protein kinase 13
 hypothetical protein FLJ11608
 v-naf musculoaponeurotic fibrosarcoma oncogene homolog F (avian)
 matrix metalloproteinase 3 (stromelysin 1, progelatinase)
 Consensus includes gb:AL137270.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434G0814 (from clone DKFZp434G0814) /FEA=mRNA /DB_XREF=gi:5807705 /UG=Hs.39 AL137270
 Consensus includes gb:N81149 /FEA=EST /DB_XREF=gi:1444476 /UG=Hs.279639 Homo sapiens mRNA; cDNA DK N81149
 carboxylate (chondroin & keratan) sulfotransferase 3
 serine/threonine kinase 23
 Consensus includes gb:AI186739 /FEA=EST /DB_XREF=gi:3737377 /DB_XREF=est:qe79c01.x1 /CLONE=IMAGE:1745184 /UG=Hs.326416 Homo sapiens mRNA; cDNA AI186739
 axonal transport of synaptic vesicles
 caldinin/calactinin-related polypeptide, alpha
 hypothetical protein FLJ11271
 pumilio (Drosophila) homolog 2
 topoisomerase (DNA) II beta (180KD)
 ATP-binding cassette, sub-family F (GCM20), member 2
 hypothetical protein
 KIAA0737 gene product
 potassium voltage-gated channel, subfamily F, member 1
 junction plakoglobin
 Consensus includes gb:AU144309 /FEA=EST /DB_XREF=gi:11005830 /DB_XREF=est:AU144309 /CLONE=HEMBA1001533 /UG=Hs.55830 Homo sapiens cDNA FLJ1001
 KIAA0217 protein
 hypothetical protein MGC10520
 CGL-53 protein
 high-mobility group (nonhistone chromosomal) protein 17-like 1
 sorting nexin 13
 cAMP responsive element binding protein-like 2
 Consensus includes gb:AI417917 /FEA=EST /DB_XREF=gi:4261421 /DB_XREF=est:ig55d04.x1 /CLONE=IMAGE:2112679 /UG=Hs.20733 Homo sapiens cDNA: FLJ22356 /AI417917
 hypothetical protein FLJ14106
 myelin basic protein
 Kruppel-associated box protein
 Consensus includes gb:BF448596 /FEA=EST /DB_XREF=gi:11514774 /DB_XREF=est:7n90b04.x1 /CLONE=IMAGE:3571902 /UG=Hs.290484 ESTs
 exportin, tRNA (nuclear export receptor for tRNAs)
 ubiquitin specific protease 2
 myosin regulatory light chain
 hypothetical protein FLJ462023.2
 KIAA0280 protein
 microtubule-associated protein lau
 Consensus includes gb:AK025352.1 /DEF=Homo sapiens cDNA: FLJ21699 fs, clone COL09829 /FEA=mRNA /DB_XREF=gi:10437851 /UG=Hs.306804 Homo sapiens cDNA: FLJ22356 /AI417917
 neuropilin 2
 chorionic somatomammotropin hormone 2
 MRE11 meiotic recombination 11 homolog A (S. cerevisiae)
 KIAA0255 gene product
 PTPN13-like, Y-linked
 phosphatase and tensin homolog (mutated in multiple advanced cancers 1)
 KIAA0808 gene product
 peptidylprolyl isomerase D (cyclophilin D)
 Consensus includes gb:AT32885 /FEA=EST /DB_XREF=gi:5053998 /DB_XREF=est:oe64d04.x5 /CLONE=IMAGE:1416391 /UG=Hs.142907 Human BRCA2 region, mRNA /AI732885
 SecE1 alpha form 2
 regulator of G-protein signalling 9
 hypothetical protein FLJ23505
 hypothetical protein FLJ13273
 FYVE and coiled-coil domain containing 1
 hypothetical protein FLJ11710
 sarcoglycan, epsilon
 lymphocyte antigen 9
 Consensus includes gb:AC006986 /DEF=Homo sapiens BAC clone RP11-155J5 from Y /FEA=CDS /DB_XREF=gi:4753246 /UG=Hs.283908 Homo sapiens BAC clone RP1AC006986
 keratin 17
 FSHD region gene 1
 actin related protein 23 complex, subunit 2 (34 kD)
 Consensus includes gb:AK022418.1 /DEF=Homo sapiens cDNA FLJ12356 fs, clone MAMMA1002339 /FEA=mRNA /DB_XREF=gi:10433811 /UG=Hs.306635 Homo sapiens cDNA: FLJ22356 /AI417917

hypothetical protein FLJ12389 similar to acetate-CoA synthetase	NM_023928	1.13
potassium channel, subfamily K, member 15 (TASK-5)	NM_022356	1.13
histidine rich calcium binding protein	NM_002152	1.13
eukaryotic translation initiation factor 4E binding protein 2	BC005057	1.13
ABO blood group (transferase A, alpha 1-3-N-acetylglucosaminyltransferase: transferase B, alpha 1-3-galactosyltransferase)	U15197	1.13
Consensus includes gb:AK024543.1 /DEF=Homo sapiens cDNA: FLJ20890 fls, clone ADKA03323. /FEA=mRNA /DB_XREF=gi:10436848 /UG=Hs.306686 Homo sapiens c AK024543	1.13	1.13
KIAA0751 gene product	NM_014677	1.13
dopamine receptor D3	NM_000796	1.13
pervin, beta	AA161053	1.13
Consensus includes gb:AA330995 /FEA=EST /DB_XREF=gi:2273701 /DB_XREF=est:n07a02.s1 /CLONE=IMAGE:985610 /UG=Hs.23729 Homo sapiens clone 24405 mRI	1.13	1.13
hypothetical protein DKFZp434E1723	AI189589	1.13
potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4	NM_002250	1.13
gb:BC001278.1 /DEF=Homo sapiens. Similar to KIAA0626 gene product, clone MGC:5129, mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0626 gene product	BC001279	1.13
Cbp/p300-interacting transactivator, with Glu/Asp-rich carboxy-terminal domain, 1	NM_004143	1.13
plakophilin 1 (ectodermal dysplasia/skin fragility syndrome)	NM_000299	1.13
synaptobin, beta 2 (dystrophin-associated protein A1, 59kD, basic component 2)	AI762021	1.13
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 6	NM_001756	1.13
ankyrin repeat and SOCS box-containing 4	AC003079	1.13
KIAA0390 gene product	NM_014717	1.13
mucosa associated lymphoid tissue lymphoma translocation gene 1	AF070528	1.13
cerebral cavernous malformations 1	NM_004912	1.13
Consensus includes gb:BE466926 /FEA=EST /DB_XREF=gi:9512701 /DB_XREF=est:h59a04.x1 /CLONE=IMAGE:3212238 /UG=Hs.279706 ESTs	BE466926	1.13
serine/threonine kinase 9	NM_003159	1.13
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 1	NM_003069	1.13
protein phosphatase methyltransferase-1	NM_016147	1.13
Rho GTPase activating protein 8	NM_017701	1.13
synaptin binding protein 1	NM_003165	1.13
adenyate kinase 2	AW277253	1.13
Consensus includes gb:R37427 /FEA=EST /DB_XREF=gi:794883 /DB_XREF=est:y55g12.s1 /CLONE=IMAGE:25997	R37427	1.13
alkaline phosphatase, placental-like 2	X07247	1.13
myosin VIIA (Usher syndrome 1B (autosomal recessive, severe))	NM_000260	1.13
CTCL tumor antigen se57-1	NM_025214	1.13
golgln-67	AI829170	1.13
chromosome 21 open reading frame 2	AI539305	1.13
voltage-dependent anion channel 3	BC002456	1.13
E2F transcription factor 6	NM_001952	1.13
vitamin D (1,25-dihydroxyvitamin D3) receptor	NM_000376	1.13
chloride channel 4	AA071195	1.13
olfactomedin 1	BC000189	1.13
interferon, alpha 10	NM_002171	1.13
cytoskeleton-associated protein 4	AW029619	1.13
hypothetical protein FLJ20132	NM_017682	1.12
fatty acid binding protein 3, muscle and heart (mammary-derived growth inhibitor)	AI041520	1.12
FLN29 gene product	NM_006700	1.12
butyrylcholinesterase	NM_000055	1.12
phosphoinositide-3-kinase, class 2, gamma polypeptide	AJ000008	1.12
cofactor required for Sp1 transcriptional activation, subunit 2 (150kD)	BE567032	1.12
poly(A) binding protein, nuclear 1	BF875004	1.12
calcitonin-related polypeptide, beta	AA747379	1.12
plasmidipin	NM_015893	1.12
olfactory receptor, family 7, subfamily E, member 13 pseudogene	AA719797	1.12
HSPCO34 protein	AV702994	1.12
Consensus includes gb:AL133262 /DEF=Human DNA sequence from clone RP1-319022 on chromosome 6 Contains part of the gene for a CDC5-like protein, the gene for	AL133262	1.12
carbonic anhydrase 1	M33987	1.12
major histocompatibility complex, class I-like sequence	AF031469	1.12
hypothetical protein MGC13098	W07700	1.12
collagen, type XIII, alpha 1	M59217	1.12
major histocompatibility complex, class II, DO beta	NM_002120	1.12
Consensus includes gb:AL523076 /FEA=EST /DB_XREF=gi:12786569 /DB_XREF=est:AL523076 /CLONE=CS00C001Y112 (3 prime) /UG=Hs.82503 H sapiens mRNA for	1.12	1.12
MRE11 meiotic recombination 11 homolog A (S. cerevisiae)	NM_005590	1.12
fragile histidine triad gene	NM_002012	1.12
PTPL1-associated RhoGAP 1	NM_004815	1.12
ribosomal protein S14	AI734124	1.12

immunoglobulin kappa constant
Consensus includes gb:AV700132 /FEA=EST /DB_XREF=gi:10302103 /DB_XREF=est:AV700132 /CLONE=GKCGSE03 /UG=Hs.295923 seven in absentia (Drosophila) nc:AV700132
U12543
KIAA0288 gene product
dnt lip and palate associated transmembrane protein 1
hypothetical protein
protease, serine, 7 (enterokinase)
hepsin (transmembrane protease, serine 1)
ATP-binding cassette, sub-family A (ABC1), member 2
bone morphogenetic protein receptor, type IA
Cas-Bc-M (murine) ectropic retroviral transforming sequence b
UDP-glucose ceramide glucosyltransferase-like 1
hypothetical protein DKFZp566133
signal sequence receptor, gamma (translocon-associated protein gamma)
RAS, member RAS oncogene family
damage-specific DNA binding protein 2 (48kD)
interleukin 9
hypothetical protein IMAGE 106914
Consensus includes gb:AL080215.1 /DEF=Homo sapiens mRNA; cDNA DKFZp588J0323 (from clone DKFZp588J0323) /FEA=mrna /DB_XREF=gi:5262706 /UG=Hs.102 AL080215
Cluster ind. A1830563;w51c03.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2406340 /clone_and=3 /gb=A1830563 /gi=5451234 /ug=Hs.120858 /len=556
hypothetical protein DKFZP434K1172
centaurin, alpha 1
solite carrier family 7 (cationic amino acid transporter, y+ system), member 2
KIAA0387 protein
amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 3
growth hormone 1
actin, alpha 3
ATPase, Na+/K+ transporting, beta 2 polypeptide
tumor necrosis factor receptor superfamily, member 8
hypothetical protein similar to tenascin-R
hypothetical protein FLJ13213
forkhead box F2
runt-related transcription factor 3
hypothetical protein MGC10646
DNA-dependent protein kinase catalytic subunit-interacting protein 2
Consensus includes gb:AJ252550 /DEF=Homo sapiens partial CK gene for glycerol kinase, exon 1 (glycerol kinase deficiency case) /FEA=CDS_1 /DB_XREF=gi:5834425 /AJ252550
KIAA0228 gene product
glutamate receptor, ionotropic, delta 2
hemochromatosis
DNA (cytosine-5-)-methyltransferase 3 alpha
NAD kinase
centrin, EF-hand protein, 1
KIAA0815 gene product
Consensus includes gb:AU146949 /FEA=EST /DB_XREF=gi:11008470 /DB_XREF=est:AU146949 /CLONE=HEM8B1001952 /UG=Hs.301817 Homo sapiens cDNA FLJ121
KIAA1056 protein
KIAA0712 gene product
solite carrier family 25 (mitochondrial carrier; Graves disease autoantigen), member 16
hydroxyprostaglandin dehydrogenase 15-(NAD)
interleukin 6 (interferon, beta 2)
cyclic AMP-regulated phosphoprotein, 21 kD
hypothetical protein FLJ20151
signal transducer and activator of transcription 1, 91kD
C-terminal binding protein 2
Kv channel-interacting protein 1
calcium channel, voltage-dependent, beta 3 subunit
UDP-GlcNAc:beta-Gal beta-1,3-N-acetylglucosaminyltransferase 6
protein phosphatase 1, regulatory subunit 3D
DKFZP434L243 protein
KIAA0790 protein
Small evolutionarily conserved RNA, resembling C/D box small nuclear
Consensus includes gb:AU147317 /FEA=EST /DB_XREF=gi:110088338 /DB_XREF=est:AU147317 /CLONE=MAMMA1000340 /UG=Hs.287491 Homo sapiens cDNA FLJ12 AU147317
synovelin
hypothetical protein FLJ10628

dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 1A	U52373	1.12
Consensus includes gb:U79273.1 /DEF=Human clone 23933 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710239 /UG=Hs.239483 Human clone 23933 mRNA sequenc	U79273	1.12
KIAA0239 protein	A1735639	1.12
hypothetical protein	AK027245	1.12
alcohol dehydrogenase 1B (class I), beta polypeptide	M24317	1.12
LW-1	NM_016153	1.12
Kruppel-like factor 12	NM_007249	1.12
ERGL protein; ERGIC-53-like protein	NM_021819	1.12
meningioma expressed antigen 5 (hyaluronidase)	AK002091	1.12
Consensus includes gb:AV661138 /FEA=EST /DB_XREF=gi:9882152 /DB_XREF=est:AV661138 /CLONE=GLCGPA11 /UG=Hs.282712 ESTs	AV661138	1.12
hypothetical protein PR01094	NM_018575	1.12
methionine adenosyltransferase 1, alpha	NM_000428	1.12
hypothetical SBB03 protein	NM_005785	1.12
proteasome gamma subfamily C, 3	BC006439	1.12
hypothetical protein FLJ20147	NM_017887	1.12
Consensus includes gb:Z69744 /DEF=H.sapiens ALL-1 gene exon 1 (and joined coding region) /FEA=CDS /DB_XREF=gi:1490270 /UG=Hs.199160 myeloidlymphoid or mit	AK025271	1.12
Consensus includes gb:AK025271.1 /DEF=Homo sapiens cDNA: FLJ21618 fs, clone COL07487. /FEA=mRNA /DB_XREF=gi:10437753 /UG=Hs.306790 Homo sapiens cD	AK025271	1.12
TAP binding protein (tapasin)	AF067286	1.12
heat shock 70kD protein 8B (mortalin-2)	A4927701	1.12
hypothetical protein FLJ21220	NM_024767	1.12
Cluster incl. AF007153:Homo sapiens clone 23763 unknown mRNA, partial cds /cds=(0,314) /gb=AF007155 /gi=2852635 /ug=Hs.186694 /len=1369	AF007155	1.12
spindlin-like	AF356353	1.12
SEC24 related gene family, member D (S. cerevisiae)	AK000709	1.12
Rho GTPase activating protein 1	BG468434	1.12
ubiquitin specific protease 7 (herpes virus-associated)	A1160440	1.12
C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 9	NM_014358	1.12
actinin	NM_006576	1.12
p53-regulated apoptosis-inducing protein 1	NM_022112	1.12
collagen, type III, alpha 1 (Ehlers-Danlos syndrome type IV, autosomal dominant)	AI813758	1.12
cystatin D	NM_001900	1.12
replication factor C (activator 1) 3 (38kD)	BC000149	1.12
cytochrome P450, subfamily IID (debrisoquine, sparteine, etc., -metabolizing), polypeptide 6	X07618	1.12
chromodomain protein, Y chromosome, 1	AF080597	1.12
muS homolog 4 (E. coli)	AF104243	1.12
cyclin D1 (PRAD1; parathyroid adenomatosis 1)	Z23022	1.12
collagen, type XVIII, alpha 1	AF018081	1.12
pre-TiNK cell associated protein	NM_021981	1.12
annexin A4	BC000182	1.12
fetal Alzheimer antigen	NM_004459	1.12
zinc finger protein 198	AL136621	1.12
N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 3	NM_004784	1.12
RNA guanylyltransferase and 5'-phosphatase	AB012143	1.12
lysosomal apyrase-like 1	NM_004901	1.12
serine/threonine kinase 17a (apoptosis-inducing)	NM_004760	1.12
ribonuclease L (2',5'-oligoadenylate synthetase-dependent)	NM_021133	1.12
serine/threonine-protein kinase PRP4 homolog	AB011108	1.12
hypothetical protein FLJ20527	NM_017863	1.12
hypothetical protein FLJ20730	NM_017945	1.12
ELK4, ETS-domain protein (SRF accessory protein 1)	NM_001973	1.12
NICE-1 protein	NM_019060	1.12
CD97 antigen	NM_001784	1.12
hypothetical protein FLJ10902	NM_018268	1.12
Consensus includes gb:AA224446 /FEA=EST /DB_XREF=gi:1844989 /DB_XREF=est:zr16a03.1 /CLONE=IMAGE:663532 /UG=Hs.311561 ESTs, Weakly similar to CYA7	AA224446	1.12
death-associated protein 6	BE965715	1.12
small nuclear RNA activating complex, polypeptide 4, 190kD	NM_003086	1.12
solute carrier family 6 (neurotransmitter transporter, GABA), member 1	A003579	1.12
hemoglobin, alpha 2	AF105974	1.12
lectin, galactoside-binding, soluble, 3 binding protein	NM_005567	1.12
cytochrome P450, subfamily IIB (phenobarbital-inducible), polypeptide 7	M29873	1.12
Consensus includes gb:AU150681 /FEA=EST /DB_XREF=gi:11012212 /DB_XREF=est:AU150681 /CLONE=NT2RP2003367 /UG=Hs.153321 Homo sapiens cDNA FLJ105	AU150681	1.12
regulator of G-protein signalling 12	AB824113	1.12
beta tubulin 1, class VI	NM_030773	1.12
hypothetical protein FLJ11565	NM_024657	1.12

KIAA0620 protein
Consensus includes gb:U80768.1 /DEF=Human EST clone 22453 mariner transposon Hsmar1 sequence. /FEA=mRNA /DB_XREF=gi:2231369 /UG=Hs.13252 Human EST U80766
Consensus includes gb:AF007147.1 /DEF=Homo sapiens clone 23712 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852625 /UG=Hs.113166 Homo sapiens clone 23712 AF007147
DKFZP564K247 protein
KIAA1031 protein
chondroectherin
organic anion transporter polypeptide-related protein 4
hypothetical protein FLJ20651
serine (or cysteine) proteinase inhibitor, clade I (neuroserpin), member 1
hypothetical protein FLJ2341
arachidonate 5-lipoxygenase
proteoglycan 3
glucagon-like peptide 1 receptor
keratin 15
nidogen (enactin)
nuclear transcription factor, X-box binding 1
cyclin-dependent kinase inhibitor 1C (p57, Kip2)
Consensus includes gb:U79300.1 /DEF=Human clone 23629 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710285 /UG=Hs.135587 Human clone 23629 mRNA sequence U79300
GDNF family receptor alpha 4
solute carrier family 22 (organic anion transporter), member 6
cytochrome P450, subfamily IVB, polypeptide 1
insulin receptor
ATPase, Na⁺/K⁺ transporting, alpha 3 polypeptide
Consensus includes gb:AK023891.1 /DEF=Homo sapiens cDNA FLJ13829 fis. clone THYRO1000825. /FEA=mRNA /DB_XREF=gi:10435964 /UG=Hs.87819 Homo sapiens, clone MGC:249; A629961
DKFZP564N1363 protein
lipin 1
prolactin receptor
mesenchymal stem cell protein DSC075
angiotensin receptor-like 1
leukocyte immunoglobulin-like receptor, subfamily A (without TM domain), member 5
Consensus includes gb:A656610 /FEA=EST /DB_XREF=gi:4740589 /DB_XREF=est:46n03.x1 /CLONE=IMAGE:2243861 /UG=Hs.87819 Homo sapiens, clone MGC:249; A629961
CD7 antigen (p41)
prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)
mucin 6, gastric
unc-51-like kinase 2 (C. elegans)
ras homolog gene family, member A
MHC class II transactivator
eosinophil chemotactic cytokine
calbindin 1, (28KD)
gamma-aminobutyric acid (GABA) A receptor, gamma 2
chromosome condensation 1-like
tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein)
BCL2-like 2
Ewing sarcoma breakpoint region 1
highly charged protein
Wolfram syndrome 1 (wolframin)
CTG26 alternate open reading frame
integrin, beta 3 (platelet glycoprotein IIb, antigen CD61)
pleckstrin homology-like domain, family A, member 1
NK-2 (Drosophila) homolog 8
hypothetical protein FLJ22054
zinc finger protein 295
hypothetical protein FLJ11535
hypothetical protein FLJ20595
H1 histone family, member 1
5-hydroxytryptamine (serotonin) receptor 4
membrane-spanning 4-domains, subfamily A, member 12 4-domains, subfamily A, member 7
myristoylated alanine-rich protein kinase C substrate
RNA-binding protein gene with multiple splicing
solute carrier family 5 (sodium/glucose cotransporter), member 1
syntaxin 1A (brain)
Wiskott-Aldrich syndrome-like
testis specific protein 1 (probe H4-1 p3-1)

aldolase B, fructose-bisphosphate
 staufen, RNA binding protein, homolog 2 (*Drosophila*)
 Consensus includes gb:AK024128.1 /DEF=Homo sapiens cDNA FLJ14066 fs, clone HEMBB1001197, /FEA=mrna /DB_XREF=gi:10436433 /UG=Hs.306665 Homo sapie
 plasminogen activator, tissue
 rRNA binding protein 1
 deafness, autosomal dominant 5
 Consensus includes gb:AF007143.1 /DEF=Homo sapiens clone 23738 mRNA sequence, /FEA=mrna /DB_XREF=gi:2852820 /UG=Hs.28205 Homo sapiens clone 23738, /AF007143
 Consensus includes gb:AF144673 /FEA=EST /DB_XREF=gi:5112883 /DB_XREF=est/tg0204.x1 /CLONE=IMAGE:2363935 /UG=Hs.186970 ESTs
 protein kinase C, beta 1
 Consensus includes gb:AA115963 /FEA=EST /DB_XREF=gi:1670996 /DB_XREF=est/tg78d11.s1 /CLONE=IMAGE:531765 /UG=Hs.323423 ESTs, Moderately similar to B AA115963
 G protein-coupled receptor 49
 basic transcription element binding protein 1
 vav 3 oncogene
 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 4
 KIAA0830 protein
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 3
 hypothetical protein FLJ20739
 v-ros UR2 sarcoma virus oncogene homolog 1 (avian)
 histone deacetylase 3
 ribosomal protein S6 kinase, 90kD, polypeptide 3
 hypothetical protein MGC4161
 neuropeptide Y receptor Y2
 aldolase B, fructose-bisphosphate
 eukaryotic translation elongation factor 1 delta (guanine nucleotide exchange protein)
 deleted in azoospermia 2
 poly(IG) binding protein 4
 hypothetical protein FLJ23053
 ACO for serine protease homologue
 plexin A2
 hypothetical protein FLJ20275
 suppressor of Ty 3 homolog (*S. cerevisiae*)
 cytochrome P450 polypeptide 43
 cholinergic receptor, nicotinic, beta polypeptide 3
 KIAA0428 gene product
 H2B histone family, member N
 hypothetical protein MGC4022
 hypothetical protein FLJ13181
 defensin, alpha 1, myeloid-related sequence
 CD86 antigen (CD28 antigen ligand 2, B7-2 antigen)
 apolipoprotein C-IV
 zinc finger protein 41
 poliovirus receptor
 COQ7 coenzyme Q, 7 homolog ubiquinone (yeast)
 polymerase (RNA) II (DNA directed) polypeptide B (140kD)
 KIAA1055 protein
 solute carrier family 25, member 13 (citrin)
 RNA binding motif protein 3
 BAI1-associated protein 1
 solute carrier family 8 (sodium/calcium exchanger), member 1
 Consensus includes gb:AK025388.1 /DEF=Homo sapiens cDNA: FLJ21735 fs, clone COLF3350, /FEA=mrna /DB_XREF=gi:10437892 /UG=Hs.306808 Homo sapiens c
 B cell RAG associated protein
 chromosome 21 open reading frame 108
 hypothetical protein FLJ23394
 Consensus includes gb:AK021636.1 /DEF=Homo sapiens cDNA FLJ11774 fs, clone HEMBA1005884, /FEA=mrna /DB_XREF=gi:10433104 /UG=Hs.293917 Homo sapie
 synixin 18
 eukaryotic translation initiation factor 3, subunit 10 (theta, 150/170kD)
 antyrim repeat, family A (RFXANK-like), 2
 KIAA0804 protein
 synovial sarcoma translocation gene on chromosome 18-like 2
 KIAA0998 protein
 syndecan 2 (heparan sulfate proteoglycan 1, cell surface-associated, fibroglycan)
 dual specificity phosphatase 2
 prostaglandin 12 (prostaglycin) synthase

Consensus includes gb:AK021987.1 /DEF=Homo sapiens cDNA FLJ11925 fs, clone HEMBS1000354. /FEA=mrna /DB_XREF=gi:10433296 /UG=Hs.191158 Homo sapie
 eiaC homolog 2 (E. coli) AF304370 1.11
 zinc finger protein 79 (p177) AA284829 1.11
 insulin-like growth factor 1 (somatomedin C) M37484 1.11
 phosphatidic acid phosphatase type 2B AV725664 1.11
 hypothetical protein FLJ20287 NM_017746 1.11
 zinc finger protein 308 BC008118 1.11
 transcription factor 7-like 2 (T-cell specific, HMG-box) AJ270770 1.11
 selectin E (endothelial adhesion molecule 1) NM_000450 1.11
 huntingtin interacting protein-1-related AB013384 1.11
 colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage) M84445 1.11
 crystallin, gamma C NM_020989 1.11
 N-ethylmaleimide-sensitive factor attachment protein, alpha BC001165 1.11
 aldehyde dehydrogenase 3 family, member A2 BC002430 1.11
 hypothetical protein from clone 24828 AF131633 1.11
 histidyl-RNA synthetase-like U18837 1.11
 ribosomal protein L10 BC003358 1.11
 regulator of G-protein signalling 12 AW170602 1.11
 Consensus includes gb:AA733079 /FEA=EST /DB_XREF=gi:2754438 /DB_XREF=est:zg79d12.s1 /CLONE=IMAGE:39957.5 /UG=Hs.286553 Homo sapiens, clone IMAGE: X82048 1.11
 WEE1+ homolog (S. pombe) NM_020119 1.11
 hypothetical protein FLB8421 BG500087 1.11
 Ras-GTPase-activating protein SH3-domain-binding protein NM_012102 1.11
 arginine-glutamic acid dipeptide (RE) repeats NM_017705 1.11
 hypothetical protein FLJ20190 NM_015652 1.11
 DKFZP564P1916 protein L36874 1.11
 synuclein, alpha (non A4 component of amyloid precursor) NM_000638 1.11
 vitronectin (serum spreading factor, somatomedin B, complement S-protein) NM_000638 1.11
 Consensus includes gb:S83374.1 /DEF=glutamate transporter II variant BH8GT IIB (5 region) human, brain and spinal cord, mRNA Partial Mutant, 129 nt. /FEA=mrna /DI S83374 1.11
 LIM homeobox protein 3 NM_006054 1.11
 Consensus includes gb:AL031178 /DEF=Human DNA sequence from clone 341E18 on chromosome 6p11.2-12.3. Contains a Serine/Threonine Protein Kinase gene (presu AF156888 1.11
 RA051-like 3 (S. cerevisiae) AL031178 1.11
 Kruppel-type zinc finger protein AF034956 1.11
 Consensus includes gb:BF594294 /FEA=EST /DB_XREF=gi:11686618 /DB_XREF=est:7h79c12.x1 /CLONE=IMAGE:3322198 /UG=Hs.287335 ESTs, Highly similar to AC1 BF594294 N21541 1.11
 DKFZP586C1620 protein AA524525 1.11
 olfactory receptor, family 2, subfamily H, member 1 AJ302604 1.11
 phospholipid scramblase 3 NM_020360 1.11
 I-complex 11 (a murine top homolog) NM_018679 1.11
 ATPase, Ca++ transporting, ubiquitous AF068220 1.11
 transcription factor (SMIF gene) NM_018403 1.11
 uracil-DNA glycosylase 2 BC004877 1.11
 major histocompatibility complex, class II, DR beta 3 AA807056 1.11
 hypothetical protein NM_019038 1.11
 suppressor of white spot homolog 2 NM_007056 1.11
 KIAA0303 protein NM_0002301 1.11
 succinate dehydrogenase complex, subunit C, integral membrane protein, 15kD D49737 1.11
 ecdonucleotide pyrophosphatase/phosphodiesterase 2 (autotaxin) L35594 1.11
 ATPase, Ca++ transporting, ubiquitous NM_005173 1.11
 myosin IIIA NM_017433 1.11
 Cluster Ind. N92708:zb44cd04.x1 Homo sapiens cDNA, 3 and /clone=IMAGE:306438 /clone_end=3 /gb=N92708 /gi=1265017 /ug=Hs.32374 /len=668 NM_001143 1.11
 amelogenin (Y chromosome) M85548 1.11
 solute carrier family 3 (cystine, dibasic and neutral amino acid transporters, activator of cystine, dibasic and neutral amino acid transport), member 1 NM_013383 1.11
 transcription factor-like 4 NM_003096 1.11
 small nuclear ribonucleoprotein polypeptide G NM_007106 1.11
 ubiquitin-like 3 BC004449 1.11
 MAD5 box transcription enhancer factor 2, polypeptide B (myocyte enhancer factor 2B) NM_000854 1.11
 glutathione S-transferase theta 2 NM_005135 1.11
 solute carrier family 12 (potassium/chloride transporters), member 6 NM_008200 1.11
 proprotein convertase subtilisin/kexin type 5 NM_002487 1.11
 Consensus includes gb:AK024867.1 /DEF=Homo sapiens cDNA: FLJ21214 fs, clone COL00523. /FEA=mrna /DB_XREF=gi:10437277 /UG=Hs.306710 Homo sapiens cDNA: AK024867 1.11
 Consensus includes gb:AK021474.1 /DEF=Homo sapiens cDNA FLJ11412 fs, clone HEMBA1000876. /FEA=mrna /DB_XREF=gi:10432668 /UG=Hs.289094 Homo sapie AK021474 1.11
 aminocyclase 1 NM_000666 1.11
 hypothetical protein DKFZp34F0318 NM_030817 1.11

muslin 5, subtypes A and C, tracheobronchial/gastric	AW192795	1.11
Williams-Beuren syndrome chromosome region 5	BC006080	1.11
guanine nucleotide binding protein (G protein), gamma transducing activity polypeptide 1	NM_021955	1.11
solute carrier family 24 (sodium/potassium/calcium exchanger), member 3	R62432	1.11
hypothetical protein FLJ12649	NM_024597	1.11
retinoblastoma 1 (including osteosarcoma)	NM_000321	1.11
cytochrome P450, subfamily IIF, polypeptide 1	NM_000774	1.11
mesoderm specific transcript homolog (mouse)	NM_002402	1.11
distal-less homeobox 4	AF254115	1.11
mal, T-cell differentiation protein	NM_002371	1.11
hypothetical protein FLJ11715	NM_024564	1.11
C-type lectin-like receptor-2	NM_016509	1.11
secure related 6 homolog (mouse)-like	AB041736	1.11
SMC1 structural maintenance of chromosomes 1-like 1 (yeast)	D80000	1.11
immunoglobulin kappa constant	BG540628	1.11
signal transducing adaptor molecule (SH3 domain and ITAM motif) 2	AI571996	1.11
dihydrolipoamide S-acyltransferase (E2 component of pyruvate dehydrogenase complex)	AW299740	1.11
hypothetical protein FLJ10932	NM_018277	1.11
basic transcription factor 3, like 1	NM_001208	1.11
Consensus includes gb:AF005487.1 /DEF=Homo sapiens MHC class II antigen (DRB6) mRNA, HLA-DRB6*0201 allele, sequence. /FEA=mRNA /DB_XREF=gi:5915893 /U AF005487	NM_022042	1.11
solute carrier family 28 (sulfate transporter), member 1	AU118165	1.11
zinc finger protein 37a (KOX 21)	X89142	1.11
keratin, hair, basic, 6 (monilehrith)	NM_005814	1.11
glycoprotein A33 (transmembrane)	AL049938	1.11
DKFZP564P1916 protein	BC005369	1.11
egl nine homolog 1 (C. elegans)	NM_014945	1.11
KIAA0843 protein	U43143	1.11
frs-related tyrosine kinase 4	NM_017555	1.11
egl nine homolog 2 (C. elegans)	AI978718	1.11
KIAA0365 gene product	NM_001921	1.11
dCMP deaminase	NM_000102	1.11
cytochrome P450, subfamily XVII (steroid 17-alpha-hydroxylase), adrenal hyperplasia	BG230586	1.11
solute carrier family 7 (cationic amino acid transporter, y+ system), member 6	NM_018274	1.11
tenasin	NM_002148	1.11
homeo box D10	AA209420	1.11
zinc finger protein 75 (D8C6)	AB006828	1.11
KIAA0280 protein	NM_003481	1.11
ubiquitin specific protease 5 (isopeptidase T)	AL042588	1.11
paternally expressed 3	AI433712	1.11
methylmalonyl Coenzyme A mutase	NM_005372	1.11
v-mos Moloney murine sarcoma viral oncogene homolog	NM_022036	1.11
G protein-coupled receptor, family C, group 5, member C	AU146004	1.11
KIAA0231 protein	AW139369	1.11
Consensus includes gb:AW295066 /FEA=EST /DB_XREF=gi:6701702 /DB_XREF=gi:6701702 /DB_XREF=est:U1-H-BW0-alt-b-09-0-U1.s1 /CLONE=IMAGE:2730209 /UG=Hs.255387 ESTs	AW295066	1.11
H2A histone family, member N	NM_003514	1.11
KIAA1816 protein	AB046836	1.11
protein phosphatase 1, regulatory (inhibitor) subunit 11	NM_021959	1.11
hypothetical protein FLJ21924	NM_024774	1.11
telomodin 1 (smooth muscle)	NM_012134	1.11
Consensus includes gb:AF052109.1 /DEF=Homo sapiens clone 23927 mRNA sequence. /FEA=mRNA /DB_XREF=gi:3360416 /UG=Hs.12473 Homo sapiens clone 23927	AF052109	1.11
myelin transcription factor 1-like	AL133024	1.11
G protein-coupled receptor 17	NM_005291	1.11
RAS2, member RAS oncogene family	AI186609	1.11
Consensus includes gb:AK023757.1 /DEF=Homo sapiens cDNA FLJ13695 fs, clone PLACE2000124. /FEA=mRNA /DB_XREF=gi:10435786 /UG=Hs.306658 Homo sapiens	AK023757	1.11
albumin	M12523	1.11
aldolase B, fructose-bisphosphate	AK026411	1.11
hypothetical protein	NM_019040	1.11
Consensus includes gb:AK000293.1 /DEF=Homo sapiens cDNA FLJ20286 fs, clone HEP04358 /FEA=mRNA /DB_XREF=gi:7020278 /UG=Hs.306391 Homo sapiens cDN	AK000293	1.11
hypothetical protein FLJ10357	R42449	1.11
gb:225434.1 /DEF=H sapiens protein-serine/threonine kinase gene, complete CDS. /FEA=mRNA /PROD=protein-serine/threonine kinase /DB_XREF=gi:405746 /FL=gb.Z25.Z25434		1.11
stromal cell-derived factor 1	NM_000809	1.11
UBX domain-containing 2	AI927512	1.11

glucagon-like peptide 1 receptor	U01157	1,11
H4 histone family, member K	NM_003546	1,11
cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 2	AF162274	1,11
CSE1 chromosome segregation 1-like (yeast)	AF053640	1,11
ectodysplasin 1, anhidrotic receptor	NM_022336	1,11
granulysin 1, anhidrotic receptor	NM_012198	1,11
granulysin, EF-hand calcium binding protein	NM_018268	1,11
hypothetical protein FLJ10904	NM_006841	1,11
transporter protein, system XI Na+ and H+-coupled glutamine transporter	AA625133	1,11
hypothetical protein FLJ2439	BC002809	1,11
down-regulator of transcription 1, TBP-binding (negative cofactor 2)	AB051447	1,11
KIAA1660 protein	Consensus includes gb:BF084105 /FEA=EST /DB_XREF=gi:10877935 /DB_XREF=est:QV4-CT0491.140900-398-e06 /UG=Hs.247954 Human thrombospondin-1 gene, pa	1,11
Consensus includes gb:BF084105 /FEA=EST /DB_XREF=gi:10877935 /DB_XREF=est:QV4-CT0491.140900-398-e06 /UG=Hs.247954 Human thrombospondin-1 gene, pa	BF084105	1,11
vacuolar protein sorting 41 (yeast)	AF135593	1,11
RNA binding motif protein BA	BG289199	1,11
v-rat musculoskeletal fibrosarcoma oncogene homolog (avian)	BF508646	1,11
RAE1 RNA export 1 homolog (S. pombe)	U85943	1,11
olfactory receptor, family 5, subfamily I, member 1	NM_006637	1,11
NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 4 (15kD, B15)	NM_004547	1,11
v-rat murine sarcoma viral oncogene homolog B1	NM_004333	1,11
ATPase, H+ transporting lysosomal (vacuolar proton pump), member M	AF077614	1,11
prostaglandin H synthase	AF010316	1,11
gb:NM_025111.1 /DEF=Homo sapiens hypothetical protein FLJ23571 (FLJ23571), mRNA, /FEA=mRNA, GEN=FLJ23571 /PROD=hypothetical protein FLJ23571 /DB_XREF=	NM_025111	1,11
phosphatidylserine decarboxylase	NM_014338	1,11
DKFZP66F0548 protein	NM_015653	1,11
nuclear receptor subfamily 2, group E, member 3	NM_014249	1,11
hypothetical protein DKFZP434A1721	NM_017609	1,11
fibrillin 2 (congenital contractual arachnodactyly)	NM_001999	1,11
Consensus includes gb:AL037534 /FEA=EST /DB_XREF=gi:5866521 /DB_XREF=est:DKFZp564F2472_1 /CLONE=DKFZp564F2472 /UG=Hs.153612 ATP-binding cassette	AL037534	1,11
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1)	AL050038	1,11
testes-specific heterogenous nuclear ribonucleoprotein G-T	NM_014469	1,11
mitochondrial ribosomal protein L20	NM_017971	1,11
sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6B	NM_020241	1,11
zinc finger protein (ZF25)	NM_016220	1,11
ATPase, Ca++ transporting, type 2C, member 1	AF189723	1,11
hypothetical protein FLJ11838	NM_024664	1,11
cytochrome P450, subfamily IVF, polypeptide 3 (leukotriene B4 omega hydroxylase)	D12620	1,11
taste receptor, type 2, member 7	NM_023919	1,11
myeloperoxidase	NM_000250	1,11
hexokinase 3 (white cell)	NM_002115	1,11
T cell receptor alpha locus	M12959	1,11
transcriptional co-activator with PDZ-binding motif (TAZ)	NM_015472	1,11
KIAA0022 gene product	NM_014880	1,11
prolyl endopeptidase	X74498	1,11
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 4	AF263541	1,11
hypothetical protein FLJ13262	NM_024914	1,11
N-acetyltransferase 1 (erythrine N-acetyltransferase)	NM_000662	1,11
interleukin 17 (cytotoxic T-lymphocyte-associated serine esterase 8)	NM_002180	1,11
KIAA0678 protein	AB014576	1,11
engulfment and cell motility 3 (ced-12 homolog, C. elegans)	NM_024712	1,11
malic enzyme 1, NADP(+)-dependent, cytosolic	NM_002395	1,11
hypothetical protein STRAAT11499	NM_021242	1,11
collagen, type IV, alpha 3 (Goodpasture antigen)	U02519	1,11
murine leukemia viral (bmt-1) oncogene homolog	NM_005180	1,11
histidine-rich glycoprotein	M13149	1,11
potassium voltage-gated channel, subfamily H (eag-related), member 2	NM_000238	1,11
apoptosis related protein	NM_014318	1,11
BCL4-like membrane protein precursor	NM_020125	1,11
hypothetical protein MGC4827	NM_024114	1,11
capicua homolog (Drosophila)	AB002304	1,11
Consensus includes gb:X07882 /DEF=Human PRB4 gene for proline-rich protein Po, allele S /FEA=mRNA /DB_XREF=gi:35647 /UG=Hs.103972 proline-rich protein BstNI : X07882		1,11
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peritonsillar biogenesis factor 12	NM_000286	1,11
Consensus includes gb:AK000173.1 /DEF=Homo sapiens cDNA FLJ20166 fls, clone COL09511, /FEA=mRNA /DB_XREF=gi:7020086 /UG=Hs.306386 Homo sapiens cDNA: AK000173		1,11

TBP-interacting protein	AB020636	1.11
breast cell glutaminase	NM_013267	1.11
hypothetical protein FLJ13386	NM_025180	1.11
angiotensin vasopressin receptor 1A	AF030625	1.11
FOS-like antigen-1	BG251266	1.11
tubor and KH domain-containing protein	NM_008862	1.11
KIAA1448 protein	H04342	1.11
gbAF13007.1 /DEF=Homo sapiens clone FLB9023 PRO2425 mRNA, complete cds. /FEA=mRNA (PROD=PRO2425 /DB_XREF=gi:11493448 /UG=Hs.302148 Homo sap AF13007.1	BF575685	1.11
karyopherin alpha 1 (importin alpha 5)	NM_005191	1.11
CD80 antigen (CD28 antigen ligand 1, B7-1 antigen)	BG477502	1.11
DEADH (Asp-Glu-Ala-Asp/His) box polypeptide 28	NM_015703	1.11
CSJ-96 protein	AL085863	1.11
dynen, axonemal, light polypeptide 4	M95528	1.11
neuron-specific protein	U25771	1.11
ADP-ribosylation factor 4-like	BC005311	1.11
hypothetical protein MGC12386	NM_018061	1.11
hypothetical protein FLJ10330	BC005934	1.11
hypothetical protein FLJ21168	AF130113	1.11
cytochrome b5 outer mitochondrial membrane precursor	M27830	1.11
M27830 Human 28S ribosomal RNA gene, complete cds (5', 3' represent transcript regions 5 prime, Middle, and 3 prime respectively)	U80756	1.11
truncaldehyde repeat containing 21	NM_006262	1.11
peripherin	NM_007223	1.11
putative G protein coupled receptor	NM_015852	1.11
Kruppel-related zinc finger protein	L10343	1.11
protease inhibitor 3, skin-derived (SKALP)	AI819115	1.11
luronate 2-sulfatase (Hunter syndrome)	NM_020355	1.11
HRPAP20 short form	NM_002531	1.11
neurotensin receptor 1 (high affinity)	NM_004644	1.11
adaptor-related protein complex 3, beta 2 subunit	NM_006692	1.11
DNA-binding protein amplifying expression of surfactant protein B	AF124790	1.11
estrogen receptor 2 (ER beta)	BC001080	1.11
hypothetical protein MGC2748	NM_004667	1.11
hect domain and RLD 2	U36501	1.11
nuclear antigen Sp100	NM_014348	1.11
similar to rat integral membrane glycoprotein POM121	AI128310	1.11
hypothetical protein FLJ21562	BC000433	1.11
Consensus includes gb:A057637 /FEA=EST /DB_XREF=gi:3331503 /DB_XREF=est:gi:3108.x1 CLONE=IMAGE:1667483 /UG=Hs.234898 ESTs, Weakly similar to 2108: A057637	AX021640	1.11
mitogen-activated protein kinase 13	AU146889	1.11
Consensus includes gb:AK021640.1 /DEF=Homo sapiens cDNA FLJ11578 fs, clone HEMBA1003571. /FEA=mRNA /DB_XREF=gi:10432861 /UG=Hs.287427 Homo sapie	AB018268	1.11
actin, alpha 2	AF016005	1.11
KIAA0725 protein	NM_021100	1.11
arginine-glutamic acid dipeptide (RE) repeats	NM_006800	1.11
NFS1 nitrogen fixation 1 (S. cerevisiae)	NM_006831	1.11
male-specific lethal 3-like 1 (Drosophila)	H05666	1.11
ATP/GTP-binding protein	NM_002986	1.11
KIAA1065 protein	U16120	1.11
small inducible cytokine subfamily D (Cys-X3-Cys), member 1 (racalactine, neurotactin)	AI806207	1.11
solute carrier family 9 (neurotransmitter transporter, taurine), member 6	NM_005890	1.11
KIAA0096 protein	NM_000805	1.11
growth arrest-specific 7	AF074000	1.11
gastrin	NM_005488	1.11
lectin, galactoside-binding, soluble, 8 (galectin 8)	U00236	1.11
target of myo1 (chicken)	NM_006380	1.11
phosphoribosyl pyrophosphate amidotransferase	NM_001512	1.11
amyloid beta precursor protein (cytoplasmic tail) binding protein 2	AL080735	1.11
glutathione S-transferase A4	BC002447	1.11
hypothetical protein DKFZp434i143	NM_004354	1.11
putative homeodomain transcription factor 1	W74442	1.11
cyclin G2	NM_003596	1.11
hypothetical protein FLJ10719	X83300	1.11
tyrosylprotein sulfotransferase 1	NM_018965	1.11
SMA4		
triggering receptor expressed on myeloid cells 2		

hypothetical protein FLJ12820	NM_022451	1.11
Consensus includes gb:AL050074.1 IDEF+Homo sapiens mRNA: cDNA DKFZ566F1946 (from clone DKFZ566F1946). /FEA=mRNA /DB_XREF=gi:4884307 /UG=Hs.27	AL050074	1.11
TATA box binding protein (TBP)-associated factor, RNA polymerase I, B, 63kD	AK026521	1.11
zinc family member 4 protein HZIC4	AF332509	1.11
retinitis pigmentosa GTPase regulator interacting protein 1	NM_020366	1.11
carbonic anhydrase IV	NM_000717	1.11
Consensus includes gb:AK022387.1 IDEF+Homo sapiens cDNA FLJ12325 fs, clone MAMMA1002125. /FEA=mRNA /DB_XREF=gi:10433772 /UG=Hs.287514 Homo sapie	AK022387	1.11
chromosome 21 open reading frame 68	NM_024944	1.11
SRY (sex determining region Y)-box 14	NM_004189	1.11
dentatorubral-pallidoluysian atrophy (atrophin-1)	D31840	1.11
SEC63 protein	AI006655	1.11
cholinergic receptor, muscarinic 3	T15991	1.11
MSTP032 protein	NM_025226	1.11
neurofilament, light polypeptide (68kD)	AL537457	1.11
Consensus includes gb:AL078636.1 IDEF+Homo sapiens mRNA (full length insert cDNA clone EUROMIMAGE 117929 /FEA=mRNA /DB_XREF=gi:5051866 /UG=Hs.227171	AL078636	1.11
paraneoplastic antigen MA2	AB020690	1.11
Consensus includes gb:U37198.1 IDEF+Homo sapiens (clone B3B3E13) Huntingtons disease candidate region mRNA fragment. /FEA=mRNA /DB_XREF=gi:600519 /UG=	U37198	1.11
RAB17, member RAS oncogene family	NM_022449	1.11
PRO1854 protein	AF180789	1.11
8-oxoguanine DNA glycosylase	NM_016620	1.11
Consensus includes gb:BF592782 /FEA=EST /DB_XREF=gi:11685106 /DB_XREF=est:7p4d06.x1 /CLONE=IMAGE:3442594 /UG=Hs.289068 Homo sapiens cDNA FLJ111	BF592782	1.11
forward box G1B	NM_005249	1.11
general transcription factor IIIC, polypeptide 4 (90kD)	NM_012204	1.11
chromosome 1 open reading frame 20	T50013	1.11
hypothetical protein HT036	NM_031207	1.11
bone morphogenetic protein 1	NM_001199	1.11
spondin 1, (I-apodin) extracellular matrix protein	AB051390	1.11
G protein-coupled receptor 35	AF089087	1.11
jerky homolog (mouse)	AF072468	1.11
odorant-binding protein 2B	NM_014581	1.11
fibroblast growth factor 3 (murine mammary tumor virus integration site (v-int-2) oncogene homolog)	NM_005247	1.11
six transmembrane epithelial antigen of the prostate	NM_012449	1.11
interleukin 1 receptor-like 1	NM_003856	1.11
chemokine (C-C motif) receptor 4	NM_005508	1.11
cathepsin Z	AF073890	1.11
maternally expressed 3	AF090934	1.11
Thy-1 cell surface antigen	AL161958	1.11
putative DNA binding protein	AL117354	1.11
Consensus includes gb:BC389837 /FEA=EST /DB_XREF=gi:13292385 /DB_XREF=est:802440589F1 /CLONE=IMAGE:4551478 /UG=Hs.289056 ESTs, highly similar to 1; BG398837	BC389837	1.10
Consensus includes gb:AW814026 /FEA=EST /DB_XREF=gi:7907020 /DB_XREF=est:RC6-ST0198-071199-021-G02 /UG=Hs.273621 Homo sapiens cDNA: FLJ21350 fs, AW814026	NM_005123	1.10
nuclear receptor subfamily 1, group H, member 4	M22921	1.10
UDP-Gal:beta-GlcNAc beta 1,4- galactosyltransferase, polypeptide 1	NM_007252	1.10
Retina-derived POU-domain factor-1	AF112345	1.10
Integrin, alpha 10	NM_022720	1.10
DiGeorge syndrome critical region gene 8	NM_005895	1.10
golgi autoantigen, golgin subfamily a, 3	NM_002138	1.10
heterogeneous nuclear ribonucleoprotein D (AU-rich element RNA binding protein 1, 37kD)	NM_007098	1.10
catrinin, heavy polypeptide-like 1	AW511135	1.10
nudix (nucleoside diphosphate linked moiety X)-type motif 4	NM_005160	1.10
adrenergic, beta, receptor kinase 2	AW134978	1.10
KIAA0984 protein	BC005111	1.10
glutamate dehydrogenase 1	NM_002781	1.10
pregnancy specific beta-1-glycoprotein 5	NM_016355	1.10
hsp256 protein	AL041124	1.10
hypothetical protein PP1665	NM_025040	1.10
hypothetical protein FLJ21841	NM_183533	1.10
Consensus includes gb:AL163533 /DEF=Human DNA sequence from clone RP11-114G1 on chromosome 13q31.3-33.1 Contains a pseudogene similar to transcription fac	AL163533	1.10
acyl-Coenzyme A oxidase 1, palmitoyl	S68189	1.10
interferon, alpha 14	NM_002172	1.10
protein inhibitor of activated STAT protein PIASy	AK022461	1.10
apoptosis regulator BCL-G	NM_030766	1.10
mitogen-activated protein kinase-activated protein kinase 2	BG325734	1.10
protein tyrosine phosphatase, receptor type, f polypeptide (PTPRF), interacting protein (liprin), alpha 3	BE501428	1.10

hypothetical protein FLJ12476
 sulfolipase, estrogen-prefering
 Consensus includes gb:AK024158.1 /DEF=Homo sapiens cDNA FLJ14098 fls, clone MAMMA100752. /FEA=mRNA /DB_XREF=gi:10436469 /UG=Hs.303289 Homo sapie
 frizzled homolog 4 (Drosophila)
 GN3 general control of amino-acid synthesis 5-like 1 (yeast)
 synaptophin, gamma 1
 Ac-like transposable element
 endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 2
 fasciculation and elongation protein zeta 2 (zyglin II)
 amyloid P component, serum
 angiotensin receptor 2
 PRO0461 protein
 hypothetical protein FLJ10520
 Cdc42 effector protein 4; binder of Rho GTPases 4
 KIAA0265 protein
 Consensus includes gb:U92706 /DEF=Human rearranged immunoglobulin heavy chain (A1VH3) gene, partial cds /FEA=CDS /DB_XREF=gi:2155271 /UG=Hs.249245 Hom
 MORF-related gene X
 hypothetical protein FLJ13105
 phosphatidylinositol glycan, class B
 Consensus includes gb:N32526 /FEA=EST /DB_XREF=gi:1152925 /DB_XREF=est|y11f04.s1 /CLONE=IMAGE:270943 /UG=Hs.300642 serologically defined colon cancer
 tumor endothelial marker 5 precursor
 endosulfine alpha
 HIRA interacting protein 5
 zinc finger protein 161 homolog (mouse)
 spectrin, beta, non-erythrocytic 2
 ephrin-A5
 hypothetical protein FLJ20085
 Consensus includes gb:AF090887.1 /DEF=Homo sapiens clone HQ0085. /FEA=mRNA /DB_XREF=gi:6690154 /UG=Hs.306562 Homo sapiens clone HQ0085
 transient receptor potential channel 4
 1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta)
 CDP-diacylglycerol synthase (phosphatidate cytidyltransferase) 2
 hypothetical protein FLJ20113
 Consensus includes gb:AK024173.1 /DEF=Homo sapiens cDNA FLJ14111 fls, clone MAMMA1001630. /FEA=mRNA /DB_XREF=gi:10436489 /UG=Hs.288567 Homo sapie
 Consensus includes gb:AK024123.1 /DEF=Homo sapiens cDNA FLJ14061 fls, clone HEMBB1000749. /FEA=mRNA /DB_XREF=gi:10436427 /UG=Hs.306664 Homo sapie
 protease, serine, 4 (trypsin 4, brain)
 chromosome 10 open reading frame 3
 C-terminal binding protein 2
 hairless homolog (mouse)
 hypothetical protein FLJ20245
 endonuclease G-like 2
 FYN oncogene related to SRC, FGR, YES
 endothelial PAS domain protein 1
 PRO1578 protein
 TAP1 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 250 kD
 gonadotropin-releasing hormone 1 (luteinizing-releasing hormone)
 cathepsin F
 DKFZP568K0524 protein
 Opa-interacting protein 5
 solute carrier family 19, member 3
 thyroid peroxidase
 carboxypeptidase E
 sodium channel, voltage-gated, type I, beta polypeptide
 origin recognition complex, subunit 6 (yeast homolog)-like
 hypothetical protein FLJ10921
 myosin IF
 T-box 2
 kinesin 1 (kinesin receptor)
 DKFZP564F1422 protein
 hypothetical protein FLJ23519
 ribosomal protein S6 kinase, 90kD, polypeptide 4
 chromosome 11 open reading frame 8
 K01391 B subtilis TrpE protein, TrpC protein corresponding to nucleotides 1883-4400 of K01391 (.5, -.M, -.3 represent transcript regions 5 prime, Middle, and 3 K01391

Consensus includes gb:AF065241.1 /DEF=Homo sapiens thioredoxin delta 3 (TXN delta 3) mRNA, partial cds. /FEA=mRNA /GEN=TXN delta 3 /PROD=thioredoxin delta 3 AF065241 1.10

Dicoforge syndrome gene H L77565 1.10

fragile X mental retardation 2 L76569 1.10

lymphoid blast crisis oncogene BF222823 1.10

ATPase, H⁺/K⁺ transporting, nongastric, alpha polypeptide NM_001676 1.10

gb:BC005914.1 /DEF=Homo sapiens, clon MGC:14510, mRNA, complete cds. /FEA=mRNA /PROD=Unknown (protein for MGC:14510) /DB_XREF=gi:13543511 /FL=gb:BC005914 1.10

CDK2-related protein kinase 7 NM_016507 1.10

Fanconi anemia, complementation group A A983201 1.10

harakuri, BCL2 interacting protein (contains only BH3 domain) U76376 1.10

insulin-like growth factor 2 receptor NM_008876 1.10

HZF-w1 protein NM_025236 1.10

tuberous sclerosis 1 AF013168 1.10

serum response factor (c-fos serum response element-binding transcription factor) A1188786 1.10

hypothetical protein FLJ20265 NM_017733 1.10

zinc finger protein 84 (HPF2) NM_003428 1.10

TNF receptor-associated factor 6 NM_004620 1.10

pleiotrophin (heparin binding growth factor 8, neurite growth-promoting factor 1) M57389 1.10

Consensus includes gb:A346181 /FEA=EST /DB_XREF=gi:4083387 /DB_XREF=est:gp49b03.x1 /CLONE=IMAGE:1928317 /UG=Hs.42712 MAX protein 1.10

DIPB protein A346181 1.10

EphA2 AA176780 1.10

hypothetical protein FLJ13195 similar to stromal antigen 3 NM_004431 1.10

carboxyterminal releasing hormone receptor 1 NM_022906 1.10

hypothetical protein FLJ13116 NM_004382 1.10

Kruppel-like factor 13 NM_016388 1.10

hypothetical protein FLJ14251 NM_015995 1.10

meningioma (disrupted in balanced translocation) 1 A614660 1.10

likely ortholog of mouse actin-related protein 8 homolog (S. cerevisiae) AA504646 1.10

retinoblastoma binding protein 2 NM_002430 1.10

chromosome 6 open reading frame 9 NM_022899 1.10

choroideremia (Rab escort protein 1) NM_005056 1.10

TAR DNA binding protein NM_000390 1.10

diethylpocamide branched chain transacylase (E2 component of branched chain keto acid dehydrogenase complex; maple syrup urine disease) BG111168 1.10

Consensus includes gb:AU146983 /FEA=EST /DB_XREF=gi:11008504 /DB_XREF=est:AU146983 /CLONE=HEMBB1002049 /UG=Hs.262212 Homo sapiens cDNA FLJ121 1.10

Homor, neuronal immediate early gene, 2 M19301 1.10

diacylglycerol kinase, epsilon (6A0D) Y19026 1.10

Bacillus subtilis /REF=M24537 /DEF=B subtilis pheB, pheA genes corresponding to nucleotides 2484-2875 of M24537 /LEN=1409 (-5, -M, -3 represent transcript regions 5' M24537 1.10

Consensus includes gb:AL021327 /DEF=Human DNA sequence from PAC 12409 on chromosome 8q21. Contains DNALJ2 (HDJ1) like pseudogene, ESTs, STSs and GSSs AL021327 1.10

ADP-ribosyltransferase (NAD⁺: poly (ADP-ribose) polymerase)-like 3 AF083068 1.10

Consensus includes gb:AV727101 /FEA=EST /DB_XREF=gi:10836522 /DB_XREF=est:AV727101 /CLONE=HTCBGF05 /UG=Hs.25272 E1A binding protein p300 /FL=gb:AV727101 1.10

HSPC059 protein NM_016536 1.10

growth factor, augmentor of liver regeneration (ERV1 homolog, S. cerevisiae) NM_005262 1.10

carboxypeptidase A4 NM_016352 1.10

sorting nexin 4 NM_003794 1.10

retinoid acid receptor responder (tazarotene induced) 2 BC000069 1.10

Consensus includes gb:AF164963.1 /DEF=Homo sapiens tumor antigen NA88-A pseudogene, complete sequence. /FEA=mRNA /DB_XREF=gi:5901726 /UG=Hs.308576 1.10

Rh type C glycoprotein NM_016321 1.10

Consensus includes gb:U86813.1 /DEF=Homo sapiens serotonin-7 receptor pseudogene, complete sequence. /FEA=mRNA /PROD=serotonin-7 receptor /DB_XREF=gi:31 U86813 1.10

insulin receptor substrate 2 BF700086 1.10

Sec23 homolog B (S. cerevisiae) BC005032 1.10

a disintegrin-like and metalloprotease (reprolysin type) with thrombospondin type 1 motif, 2 NM_014244 1.10

putative secreted protein NM_015926 1.10

T-cell lymphoma invasion and metastasis 1 NM_003253 1.10

Consensus includes gb:AF195953 /DEF=Homo sapiens membrane-bound aminopeptidase P (XNPEP2) gene, complete cds. /FEA=mRNA /DB_XREF=gi:11066156 /UG=Hs. AF039555 1.10

visinin-like 1 NM_013332 1.10

hypoxia-inducible protein 2 NM_017996 1.10

hypothetical protein FLJ10103 NM_022844 1.10

chorionic somatomammotropin hormone 2 AA129726 1.10

HYA22 protein U18759 1.10

nuclear factor I/X (CCAAT-binding transcription factor) NM_016578 1.10

HBV pX associated protein-8 NM_008865 1.10

KUAI069 protein AW665865 1.10

Consensus includes gb:AW971983 /FEA=EST /DB_XREF=gi:8161949 /DB_XREF=est:EST384192 /UG=Hs.293003 ESTs, Moderately similar to ALU4_HUMAN ALU SUBF AW971983 1.10

NGF1A binding protein 2 (EGR1 binding protein 2)
 gb:NM_024907.1 /DEF=Homo sapiens hypothetical protein FLJ11798 (FLJ11798), mRNA, /FEA=mRNA /GEN=FLJ11798 /PROD=hypothetical protein FLJ11798 /DB_XREF=NM_024907
 hypothetical protein FLJ22529
 yeast-associated membrane protein 1
 neuropilin 1
 immunoglobulin kappa variable 1D-8
 zinc finger protein 278
 N-myristoyltransferase 1
 ubiquitin specific protease 22
 DESC1 protein
 phosphatase, glycogen; liver (Hers disease, glycogen storage disease type VI)
 chromosome 22 open reading frame 4
 human T-cell leukemia virus enhancer factor
 nucleoside diphosphate kinase type 6 (inhibitor of p53-induced apoptosis-alpha)
 chloride channel 5 (nephrolithiasis 2, X-linked; Dent disease)
 KIAA0871 protein
 Chediak-Higashi syndrome 1
 transcription factor CP2
 progastrin (pepsinogen C)
 neurofibromin 2 (bilateral acoustic neuroma)
 methyl CpG binding protein 2 (Rett syndrome)
 cytochrome P450, 51 (lanosterol 14-alpha-demethylase)
 hypothetical protein FLJ10420
 septaplerin reductase (7,8-dihydrobiopterin:NADP+ oxidoreductase)
 deiodinase, iodothyronine, type II
 splicing factor (45KD)
 sec13-like protein
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 9
 general transcription factor IIE, polypeptide 1 (alpha subunit, 56kD)
 hypothetical protein MGC11061
 forkhead box D1
 KIAA0869 protein
 RNA binding motif protein 7
 slug homolog, zinc finger protein (chicken)
 mitochondrial ribosomal protein 63
 cyclin-dependent kinase 5, regulatory subunit 2 (p39)
 KIAA0749 protein
 hypothetical protein FLJ20628
 translin-associated factor X
 membrane component, chromosome 17, surface marker 2 (ovarian carcinoma antigen CA125)
 prostaglandin I2 (prostaglandin) receptor (IP)
 hypothetical protein FLJ23306
 Consensus includes gb:AI093572 /FEA=EST /DB_XREF=gi:3432548 /DB_XREF=est:gb15111.x1 /CLONE=IMAGE:1696365 /UG=Hs, 18081 Homo sapiens clone DT1P186
 mannosyl (beta-1,4)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase
 hypothetical protein FLJ12875
 stratifin
 ubiquitin-conjugating enzyme E2L3
 hypothetical protein FLJ13659
 Consensus includes gb:AL109709.2 /DEF=Homo sapiens mRNA full length insert cDNA clone EUROIMAGE 43432 /FEA=mRNA /DB_XREF=gi:187596 /UG=Hs, 167456
 diacylglycerol kinase, gamma (90kD)
 synovial sarcoma, X breakpoint 5
 DKFZP564D116 protein
 Sp2 transcription factor
 cell division cycle 42 (GTP binding protein, 25kD)
 protein phosphatase, EF hand calcium-binding domain 1
 hypothetical protein FLJ21069
 growth arrest-specific 11
 adaptor-related protein complex 4, epsilon 1 subunit
 ribosomal protein L28
 DAZ associated protein 2
 YY1-associated factor 2
 chloride channel 4
 mitogen-activated protein kinase kinase 4

capping protein (actin filament) muscle Z-line, alpha 1
 pregnancy-associated plasma protein-E
 protease, serine, 11 (IGF binding)
 Cdc42 effector protein 3
 Fanconi anemia, complementation group A
 hypothetical protein FLJ10850
 PAP-1 binding protein
 synaptogyrin 4
 KIAA0972 protein
 arginine vasopressin receptor 1A
 DKFZP564O0423 protein
 REV3-like, catalytic subunit of DNA polymerase zeta (yeast)
 trinucleotide repeat containing 1
 annexin A11
 signal transducer and activator of transcription 1, 91kD
 zinc finger protein 278
 CDC14 cell division cycle 14 homolog A (S. cerevisiae)
 neutralized-like (Drosophila)
 Consensus includes gb:BE794663 /FEA=EST /DB_XREF=gi:10215861 /DB_XREF=est:601590459F1 /CLONE=IMAGE:3944463 /UG=Hs.306668 Homo sapiens cDNA FLJ BE794663
 GLI-Kruppel family member GLI2
 gamma-glutamyl hydrolase (conjugase, poly(glycoprotein)glutaminyl hydrolase)
 interleukin 6 signal transducer (gp130, oncostatin M receptor)
 glycine N-methyltransferase
 Consensus includes gb:AK022254.1 /DEF=Homo sapiens cDNA FLJ12192 lis, clone MAMMA1000851, /FEA=mRNA /DB_XREF=gi:10433612 /UG=Hs.288451 Homo sapiens cDNA FLJ BE794663
 complement component 1, q subcomponent, receptor 1
 paired mesoderm homeobox 2b
 Cluster /ind: A1472320 /j87c02.x1 Homo sapiens cDNA, 3 end /clone=IMAGE:2148482 /clone_end=3 /gb=A1472320 /gi=4334410 /ug=Hs.48504 /len=548
 smoothelin
 C-terminal binding protein 2
 PRO0297 protein
 centaurin, delta 2
 hypothetical protein DKFZp586E1923
 myocilin, trabecular meshwork inducible glucocorticoid response
 hypothetical protein FLJ14220
 Consensus includes gb:AL035308.1 /DEF=H. sapiens gene from PAC 42816, /FEA=CDS /PROD=hypothetical protein /DB_XREF=gi:4200237 /UG=Hs.105911 H. sapiens ge
 homolog of mouse quaking QKI (KH domain RNA binding protein)
 ankrym 3, node of Ranvier (ankyrin G)
 hypothetical protein FLJ10210
 dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2
 natriuretic peptide receptor Aguanalyse cyclase A (atrionatriuretic peptide receptor A)
 aminolevulinatase, delta-, dehydratase
 core-binding factor, runt domain, alpha subunit 2, translocated to, 2
 FYN oncogene related to SRC, FGR, YES
 hypothetical protein FLJ2242
 neuronal protein
 hypothetical protein FLJ12455
 Consensus includes gb:AW469184 /FEA=EST /DB_XREF=gi:7039280 /DB_XREF=est:hc78p04.x1 /CLONE=IMAGE:2898870 /UG=Hs.65408 ESTs
 Consensus includes gb:AL137325.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M0835 (from clone DKFZp434M0835), /FEA=mRNA /DB_XREF=gi:6807816 /UG=Hs.27
 small inducible cytokine subfamily A (Cys-Cys), member 15
 crystallin, beta B3
 galanin-related peptide
 aldehyde dehydrogenase 7 family, member A1
 chromodomain helicase DNA binding protein 2
 Consensus includes gb:U05589.1 /DEF=Human ribosomal protein S1 homolog mRNA, partial cds, /FEA=mRNA /DB_XREF=gi:497001 /UG=Hs.371 ribosomal protein S1-III U05589
 gb:U72513.1 /DEF=Human RPL13-2 pseudogene mRNA, complete cds, /FEA=mRNA /PROD=RPL13-2 /DB_XREF=gi:1673516 /UG=Hs.159486 Human RPL13-2 pseudog
 Kruppel-like factor 5 (intestinal)
 nuclear protein, axia-teleangiectasia locus
 achaeate-scut complex-like 1 (Drosophila)
 POU domain, class 5, transcription factor 1
 CD1E antigen, e polypeptide
 zinc finger protein 192
 collagen-like tail subunit (single strand of homotrimer) of asymmetric acetylcholinesterase
 Consensus includes gb:A1279819 /FEA=EST /DB_XREF=gi:3918053 /DB_XREF=est:qm26h04.x1 /CLONE=IMAGE:1882999 /UG=Hs.27373 Homo sapiens mRNA; cDNA [A1279819

putative DNA/chromatin binding motif
 Consensus includes gb:AK024142.1 /DEF=Homo sapiens cDNA FLJ14080 fls, clone HEMBB1002152. /FEA=mRNA /DB_XREF=gi:10436451 /UG=Hs.301905 Homo sapiens
 hypothetical protein FLJ21172
 hypothetical protein FLJ22457
 putative DNA binding protein
 spastic paraplegia 4 (autosomal dominant; spastin)
 KIA00562 gene product
 Consensus includes gb:AU145025 /FEA=EST /DB_XREF=gi:11006546 /DB_XREF=est:AU145025 /CLONE=HEMBA1003667 /UG=Hs.296748 Homo sapiens cDNA FLJ14141 /AU145025
 talin 2
 ATPase, Class II, type 9B
 protocadherin gamma subfamily C, 3
 ribosomal protein L5
 Consensus includes gb:AK024909.1 /DEF=Homo sapiens cDNA: FLJ21256 fls, clone COL01402. /FEA=mRNA /DB_XREF=gi:10437324 /UG=Hs.306724 Homo sapiens cDNA
 DKFZP564O123 protein
 mitogen inducible 2
 KIA00887 protein
 selectin P ligand
 Consensus includes gb:W26966 /FEA=EST /DB_XREF=gi:1306194 /DB_XREF=est:16h5 /UG=Hs.293838 ESTs
 cytochrome P450 monooxygenase
 synovial sarcoma, X breakpoint 2
 orosomucoid 1
 hypothetical protein MGC5487
 prostaglandin E receptor 3 (subtype EP3)
 wingless-type MMTV integration site family, member 6
 paired mesoderm homeo box 1
 nuclear pore complex protein
 hypothetical protein FLJ00001
 X17013 B subunit Iys gene for diaminopimelate decarboxylase corresponding to nucleotides 350-1345 of X17013 (-5, -M, -3 represent transcript regions 5 prime, Middle, ar
 autocrine motility factor receptor
 chorionic somatomammotropin hormone-like 1
 ring finger protein (C3HC4 type) 8
 growth hormone releasing hormone receptor
 zinc finger protein 264
 PBX/notch1 1 homeobox 2
 nuclear antigen Sp100
 Consensus includes gb:AL031737 /DEF=Human DNA sequence from clone 8B22 on chromosome 1p35.1:36.21 Contains gene similar to cytoplasmic dynein light chain 1, (AL031737)
 hypothetical protein LOC57333
 Friedreich ataxia region gene X123
 ubiquitin-conjugating enzyme E21 (UBC9 homolog, yeast)
 collagen, type XIV, alpha 1 (undulin)
 synuclein, alpha (non A4 component of amyloid precursor)
 DKFZP566A0522 protein
 ectonucleotide pyrophosphatase/phosphodiesterase 3
 hypothetical protein
 cell division cycle 27
 hypothetical protein MGC10791
 Cdo42 effector protein 4; binder of Rho GTPases 4
 biglycan
 KIAA0930 protein
 crystallin, beta B2
 Ric-like, expressed in many tissues (Drosophila)
 hypothetical protein PRO1318
 sterile alpha and HEAT/Armadillo motif protein, ortholog of Drosophila
 nudix (nucleoside diphosphate linked moiety X)-type motif 3
 CD24 antigen (small cell lung carcinoma cluster 4 antigen)
 uncharacterized hypothalamus protein HT008
 5-hydroxytryptamine (serotonin) receptor 1A
 Melanoma associated gene
 solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1
 LIM domain protein
 protein tyrosine phosphatase, non-receptor type 14
 clock homolog (mouse)
 hematopoietically expressed homeobox

corticotropin releasing hormone binding protein NM_001882 1.09
 Consensus includes gb:AL122093.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434B2115 (from clone DKFZp434B2115), /FEA=mRNA /DB_XREF=gi:6102904 /UG=Hs.13 /AL122093 1.09
 guanylate cyclase 2D, membrane (retina-specific) NM_000180 1.09
 eukaryotic translation initiation factor 4E binding protein 2 AW500473 1.09
 small inducible cytokine subfamily A (Cys-Cys), member 8 (monocyte chemotactic protein 2) A184980 1.09
 corticotropin releasing hormone receptor 2 NM_001883 1.09
 contactin associated protein 1 NM_003632 1.09
 pili-like transcription factor NM_012228 1.09
 AF209929 1.09
 NM_004199 1.09
 proadrenomedullin-2, 2-oxoglutarate 4-dioxygenase (proline 4-hydroxylase), alpha polypeptide II T15766 1.09
 Consensus includes gb:T15766 /FEA=EST /DB_XREF=gi:517828 /DB_XREF=est:181859 /UG=Hs.321572 Homo sapiens clone 2504.8 mRNA sequence NM_001483 1.09
 glioblastoma amplified sequence NM_001118 1.09
 adenylate cyclase activating polypeptide 1 (pituitary) receptor type I BC006230 1.09
 monoglyceride lipase NM_004428 1.09
 ephrin-A1 NM_013989 1.09
 transition initiation factor IF2 AB018284 1.09
 C3HC4-like zinc finger protein NM_016422 1.09
 KIAA1067 protein AK023632 1.09
 PRKC, apoptosis, WT1, regulator A1780470 1.09
 5-hydroxytryptamine (serotonin) receptor 2B NM_000867 1.09
 peroxisomal biogenesis factor 11A NM_003847 1.09
 zinc finger protein 272 X78931 1.09
 gb:AF049656.1 /DEF=Homo sapiens inducible nitric oxide synthase (NOS) mRNA, complete cds. /FEA=CDS /GEN=INOS /PROD=inducible nitric oxide synthase /DB_XREF AF049656 1.09
 Cluster incl. A1689655, wc12005.x1 Homo sapiens cDNA, 3' end /done=IMAGE:2314928 /done_end=3 /gb=A1689655 /gi=4834428 /ug=Hs.30251 /len=649 A1689655 1.09
 Consensus includes gb:AL121948 /DEF=Human DNA sequence from clone RP3-398E4 on chromosome 6. Contains the 3' end of the SIM1 gene for single-minded (Drosop AL121948 1.09
 Consensus includes gb:BCG536224 /FEA=EST /DB_XREF=gi:13527769 /DB_XREF=est:60256445F1 /CLONE=IMAGE:4690258 /UG=Hs.123030 Human kappa-immunoglobulin BG536224 1.09
 gb:AF090900.1 /DEF=Homo sapiens clone HQ0189 PRO0189 mRNA, complete cds. /FEA=mRNA /PROD=PRO0189 /DB_XREF=gi:6690176 /UG=Hs.91393 Homo sapiens: AF090900 W22117 1.09
 HT1017 protein NM_017831 1.09
 hypothetical protein FLJ20699 NM_004720 1.09
 endothelial differentiation, lysophosphatidic acid G-protein-coupled receptor, 4 NM_022470 1.09
 p53 target zinc finger protein NM_001091 1.09
 amiloride binding protein 1 (amine oxidase (copper-containing)) BC005297 1.09
 kynurenine 3-monooxygenase (kynurenine 3-hydroxylase) NM_004619 1.09
 TNF receptor-associated factor 5 NM_019598 1.09
 gb:NM_019598.1 /DEF=Homo sapiens kallikrein 12 (KLK12), mRNA. /FEA=mRNA /GEN=KLK12 /PROD=kallikrein 12 /DB_XREF=gi:9665235 /UG=Hs.159679 kallikrein 12 NM_019598 1.09
 nuclear receptor subfamily 2, group F, member 2 AV703465 1.09
 hypothetical protein PRO1331 NM_030778 1.09
 putative Rab5 GDP/GTP exchange factor homologue NM_017801 1.09
 hypothetical protein FLJ20396 NM_014504 1.09
 KIAA0995 protein AB023212 1.09
 carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 5 NM_012126 1.09
 protodactherin 12 BF511556 1.09
 peptidylprolyl isomerase E (cyclophilin E) NM_006738 1.09
 lymphoid blast crisis oncogene A1701514 1.09
 claudin 9 AV756161 1.09
 hypothetical protein FLJ22059 BF871416 1.09
 DKFZP386N0721 protein BF575588 1.09
 myeloid/lymphoid or mixed-lineage leukemia (lithorax homolog, Drosophila), translocated to, 2 NM_004945 1.09
 dynamin 2 BC004917 1.09
 hypothetical protein MGC4771 NM_018166 1.09
 hypothetical protein FLJ10647 BG287153 1.09
 mannosidase, alpha, class 1A, member 1 AF072826 1.09
 ras responsive element binding protein 1 BC000249 1.09
 phosphodiesterase 6B, cGMP-specific, rod, beta (congenital stationary night blindness 3, autosomal dominant) NM_000614 1.09
 ciliary neurotrophic factor NM_017770 1.09
 elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 2 BC006141 1.09
 erythrocyte membrane protein band 4.1-like 3 AF113514 1.09
 histone acetyltransferase NM_017423 1.09
 UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (GalNAc-T7) NM_001234 1.09
 caveolin 3 D32047 1.09
 thrombospondin (myeloproliferative leukemia virus oncogene ligand, megakaryocyte growth and development factor) NM_018413 1.09
 chondroitin 4-sulfotransferase

T cell receptor alpha locus	AE000659	1.09
phosphatidylinositol glycan, class L	AI742901	1.09
slit homolog 2 (Drosophila)	AF055585	1.09
hypothetical protein FLJ22393	NM_025106	1.09
heat shock 90kD protein 1, alpha	BG420237	1.09
hypothetical protein FKSG28	W28828	1.09
deoxyribonuclease I	M55983	1.09
glycerol kinase	X88285	1.09
UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 6	NM_004775	1.09
mutL homolog 3 (E. coli)	NM_014381	1.09
sterile alpha and HEAT/Armadillo motif protein, ortholog of Drosophila	AJ280445	1.09
Consensus includes gb:AW593996 /FEA=EST /DB_XREF=gi:7281254 /DB_XREF=est:hg41g06.x1 /CLONE=IMAGE:2948218 /UG=Hs:124969 Homo sapiens clone 24707	AW593996	1.09
KIAA0687 protein	AI978623	1.09
KIAA0435 gene product	AB007895	1.09
KIAA0674 protein	AB014574	1.09
Consensus includes gb:AF090883.1 /DEF=Homo sapiens clone HQ0021. /FEA=mRNA /DB_XREF=gi:6690149 /UG=Hs:306560 Homo sapiens clone HQ0021	AF090883	1.09
nasopharyngeal carcinoma susceptibility protein	NM_013275	1.09
DKFZP564B167 protein	NM_015415	1.09
coronadomesin	NM_001284	1.09
MORF-related gene 15	NM_006791	1.09
Consensus includes gb:AL031600 /DEF=Human DNA sequence from clone 390E6 on chromosome 18. Contains the 3 part of the CLCN7 gene for chloride channel 7. EST: LQ3380	NM_003160	1.09
gonadotropin-releasing hormone receptor	AL080137	1.09
DKFZP434J193 protein	NM_012101	1.09
tripartite motif-containing 28	AI984421	1.09
GpE-like protein co-chaperone	AF229062	1.09
death effector filament-forming Ced-4-like apoptosis protein	NM_017612	1.09
hypothetical protein DKFZP434E2220	AF229062	1.09
Consensus includes gb:AL049449.1 /DEF=Homo sapiens mRNA: cDNA DKFZP586B1722 (from clone DKFZP586B1722) /FEA=mRNA /DB_XREF=gi:4500234 /UG=Hs:80	AL049449	1.09
KIAA0805 protein	AI815033	1.09
centromere protein F (350/400kD, mitotin)	U30872	1.09
nucleolar protein 1 (120kD)	NM_006170	1.09
chromatin assembly factor 1, subunit A (p150)	BF000239	1.09
ephin-B3	NM_001406	1.09
brain-specific angiogenesis inhibitor 2	NM_001703	1.09
inhibitor of Brulon's tyrosine kinase	AF235049	1.09
ectodermal-neural crest (with BTB-like domain)	NM_003633	1.09
PRO0811 protein	NM_014076	1.09
insulin-like growth factor 2 (somatomedin A)	M17663	1.09
oculocutaneous albinism II (pink-eye dilution (murine) homolog)	NM_000275	1.09
core promoter element binding protein	U51869	1.09
perilipin	NM_002666	1.09
alpha-methylacyl-CoA racemase	AA88589	1.09
keratin, hair, basic, 4	Y19209	1.09
Consensus includes gb:AL137715.1 /DEF=Homo sapiens mRNA: cDNA DKFZP434K0423 (from clone DKFZP434K0423); partial cds. /FEA=mRNA /GEN=DKFZP434K0423	AL137715	1.09
ribosomal protein L21	AL356414	1.09
FtsJ homolog 1 (E. coli)	NM_012280	1.09
extracellular matrix protein 1	U65932	1.09
microtubule-associated protein like echinoderm EMAP	NM_012155	1.09
cell division cycle 25A	AI343459	1.09
hypothetical protein FLJ20298	NM_017752	1.09
amyloid beta (A4) precursor protein-binding, family A, member 2 (X11-like)	AB014719	1.09
proline-serine-threonine phosphatase interacting protein 2	NM_024430	1.09
alpha-fetoprotein	NM_001134	1.09
transcription factor AP-2 alpha (activating enhancer binding protein 2 alpha)	NM_003220	1.09
RAD52 homolog (S. cerevisiae)	AF125949	1.09
proline-rich protein BstNI subfamily 1	NM_005039	1.09
casepase 4, apoptosis-related cysteine protease	U25804	1.09
calycon, D1 dopamine receptor-interacting protein	NM_015722	1.09
folistatin-like 3 (secreted glycoprotein)	NM_005880	1.09
kinesin-like 1	NM_004523	1.09
CGI-111 protein	NM_016048	1.09
Consensus includes gb:AC004490 /DEF=Homo sapiens chromosome 19, cosmid R29381 /FEA=CDS /DB_XREF=gi:2995576 /UG=Hs:156015 Homo sapiens chromosome AC004490	AC004490	1.09
chromosome 1 open reading frame 22	NM_025191	1.09

DKF2P564M082 protein
 intracisternal A particle-promoted polypeptide
 ribosomal protein L17
 DNA-crosslink repair gene SNM1
 acetyl-Coenzyme A carboxylase beta
 ubiquitin-conjugating enzyme E2B (RAD6 homolog)
 sedmin-like 2
 regulatory factor X-associated protein
 Consensus includes gb:U79248.1 /DEF=Human clone 23826 mRNA sequence, /FEA=mRNA /DB_XREF=gi:170195 /UG=Hs.12484 Human clone 23826 mRNA sequence
 diaphanous homolog 3 (Drosophila)
 delta-tubulin
 HCGIV-6 protein
 Consensus includes gb:AJ307760 /FEA=EST /DB_XREF=gi:4002364 /DB_XREF=eat:ib24p09.x1 /CLONE=IMAGE:2055328 /UG=Hs.14074 Homo sapiens clone 24627 mR
 S100 calcium binding protein A8 (calgranulin A)
 glycoporphin A (includes MN blood group)
 TAF-like RNA polymerase II, p300/CBP-associated factor (PCAF)-associated factor, 65 kD
 hypothetical protein FLJ20281
 Consensus includes gb:AW341200 /FEA=EST /DB_XREF=gi:6837826 /DB_XREF=est:xx295g12.x1 /CLONE=IMAGE:2872006 /UG=Hs.188882 Homo sapiens cDNA: FLJ21
 adaptor-related protein complex 3, delta 1 subunit
 UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter
 serine palmitoyltransferase, long chain base subunit 2
 serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
 claudin 15
 phospholipase C, beta 2
 cholinergic receptor, nicotinic, alpha polypeptide 6
 myeloid leukemia factor 1
 plasmalemma vesicle associated protein
 H328
 isovaleryl Coenzyme A dehydrogenase
 regulatory factor X, 3 (influences HLA class II expression)
 homolog of yeast MOG1
 Consensus includes gb:AK026238.1 /DEF=Homo sapiens cDNA: FLJ22585 fa, clone HSI02678, /FEA=mRNA /DB_XREF=gi:10439027 /UG=Hs.287701 Homo sapiens cD
 Consensus includes gb:AL136318 /DEF=Human DNA sequence from clone RP1-254P11 on chromosome 10 Contains ESTs, STSs and GSSs. Contains a gene for a novel
 adducin 1 (alpha)
 protocadherin 11
 nucleosome assembly protein 1-like 2
 T cell receptor alpha locus
 RCD1 required for cell differentiation1 homolog (S. pombe)
 protein x 0004
 prostate epithelium-specific Ets transcription factor
 chondroitin sulfate proteoglycan 2 (versican)
 inhibitor of DNA binding 4, dominant negative helix-loop-helix protein
 acrosomal vesicle protein 1
 hypothetical protein FLJ22686
 low density lipoprotein receptor (familial hypercholesterolemia)
 selectin P (granule membrane protein 140kD, antigen CD62)
 hypothetical protein FLJ20280
 ficollin (collagen/vitronectin domain-containing) 3 (Hakata antigen)
 interferon regulatory factor 5
 death effector filament-forming Ced-4-like apoptosis protein
 U14573 Human Alu-Sq subfamily consensus sequence.
 related RAS viral (-ras) oncogene homolog
 terasin
 Rho guanine nucleotide exchange factor (GEF) 3
 ras-related C3 botulinum toxin substrate 1 (rho family, small GTP binding protein Rac1)
 microtubule-associated protein 7
 DKF2P34O047 protein
 decorin
 chromosome 16 open reading frame 7
 KIAA0825 protein
 neuropilin 1
 hypothetical protein PRO0388
 Consensus includes gb:AU147598 /FEA=EST /DB_XREF=gi:11009119 /DB_XREF=est:AU147598 /CLONE=MAMMA1001143 /UG=Hs.296707 Homo sapiens cDNA FLJ12 AU147598

NM_014042
 NM_005897
 AL049597
 D42045
 R99037
 A4877765
 NM_007029
 NM_000538
 NM_017748
 U78248
 NM_030932
 BC000258
 AF036973
 NM_002964
 A1240545
 BF058726
 NM_017748
 NM_01341200
 A1424923
 N80922
 U15555
 AF19873
 NM_014343
 BC000939
 NM_004198
 NM_022443
 AF326591
 NM_015726
 M34192
 NM_002919
 NM_014185
 AK026238
 AL136318
 AL556041
 NM_014522
 NM_021963
 AW508865
 A1567462
 NM_016301
 A1745526
 BF590263
 AL022726
 NM_020114
 NM_024957
 A861942
 NM_003005
 NM_017741
 NM_003665
 A1028035
 AF229061
 U14573
 NM_006270
 AF116810
 NM_019535
 BC004247
 AJ242502
 NM_015594
 AF138302
 NM_004913
 NM_015046
 NM_013957
 NM_018562
 NM_018562

microtubule-associated protein, RP/EB family, member 2	AB016823	1.09
goosecoid-like	NM_005315	1.09
hypothetical protein FLJ23468	NM_024629	1.09
KIAA1026 protein	AB015329	1.09
KIAA1088 protein	AB029012	1.09
LentC (bacterial) antibiotic synthetase component C)-like 1	AI935255	1.09
hypothetical protein FLJ21551	NM_024801	1.09
hypothetical protein FLJ10110	NM_017998	1.09
Machado-Joseph disease (spinocerebellar ataxia 3, olivopontocerebellar ataxia 3, autosomal dominant, ataxin 3)	AI686099	1.09
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 4 (GalNAc-T4)	NM_003774	1.09
Siegrin syndrome antigen A2 (60kD, ribonucleoprotein autoantigen SSA/Ro)	AL538601	1.09
hypothetical protein FLJ21140	NM_024778	1.09
proteasome (prosome, macropain) 26S subunit, non-ATPase, 9	BC002383	1.09
zinc finger protein 274	NM_016325	1.09
hypothetical protein FLJ20241	AW163030	1.09
carboxypeptidase N, polypeptide 1, 50kD	NM_001941	1.09
desmocollin 3	M32221	1.09
RNA isopentenylpyrophosphate transferase	NM_017846	1.09
ras inhibitor	NM_004292	1.09
olfactory receptor, family 2, subfamily F, member 1	NM_012369	1.09
Consensus includes gb:L23852.1 /DEF=Homo sapiens (clone Z146) retinal mRNA, 3' end and repeat region /FEA=mRNA /DB_XREF=gi:393126 /UG=Hs.73838 Homo sapiens	NM_006948	1.09
stress 70 protein chaperone, microsome-associated, 60kD	L12468	1.09
glutaryl aminopeptidase (aminopeptidase A)	NM_004216	1.09
death effector domain-containing	NM_004172	1.09
solute carrier family 1 (glial high affinity glutamate transporter), member 3	NM_012072	1.09
complement component 1, q subcomponent, receptor 1	NM_018014	1.09
B-cell CLL/lymphoma 11A (zinc finger protein)	NM_024025	1.09
hypothetical protein MGC1136	BE542684	1.09
eukaryotic translation initiation factor 1A	NM_005971	1.09
FXYD domain-containing ion transport regulator 3	NM_006185	1.09
nuclear riboflavin synthase protein 1	BC006163	1.09
dynactin 1 (p150, glued homolog, Drosophila)	NM_015925	1.09
liver-specific bHLH-Zip transcription factor	NM_020189	1.09
OC6 protein	AB051446	1.09
KIAA1639 protein	NM_024671	1.09
hypothetical protein FLJ23438	L23982	1.09
collagen, type VII, alpha 1 (epidermolysis bullosa, dystrophic, dominant and recessive)	AI432713	1.09
Consensus includes gb:AI432713 /FEA=EST /DB_XREF=gi:4283899 /DB_XREF=est:th3e02.x1 /CLONE=IMAGE:2121050 /UG=Hs.241053 ESTs	M99578	1.09
DNA segment on chromosome X and Y (unique) 155 expressed sequence	U73304	1.09
cannabinoid receptor 1 (brain)	AI650257	1.09
alpha thalassemia/mental retardation syndrome X-linked (RAO54 homolog, S. cerevisiae)	AI479175	1.09
lipoprotein lipase	BF872975	1.09
H2.O-like homeo box 1 (Drosophila)	M60721	1.09
mannosidase, alpha, class 1A, member 2	H87840	1.09
likely ortholog of mouse zinc finger homeodomain 4	NM_024721	1.09
regulated in glioma	NM_006394	1.09
GDP-mannose 4,6-dehydratase	AI762113	1.09
hypothetical protein FLJ13322	NM_024722	1.09
nuclear receptor coactivator 3	AI761748	1.09
adenylyl cyclase-associated protein 2	NM_006366	1.09
hypothetical protein MGC4707	NM_024113	1.09
KIAA0675 gene product	BG502305	1.09
KIAA0738 gene product	BC000609	1.09
zinc finger protein 151 (pH2-87)	AW574837	1.09
cytochrome P450, subfamily IVF, polypeptide 11	NM_021187	1.09
heat shock 70kD protein 6 (HSP70B)	X51757	1.09
KIAA0185 protein	AI547263	1.09
SEX gene	AI675453	1.09
protein tyrosine phosphatase, receptor type, U	U71075	1.09
Ras-induced senescence 1	BF062629	1.09
endonuclease G-like 1	BF511976	1.09

hypothetical protein FLJ20170	NM_017696	1.09
Consensus includes gb:BF155672 /FEA=EST /DB_XREF=gi:11050855 /DB_XREF=est:QV4-HT0894-290900-399-c04 /UG=Hs.146200 ESTs	BF155672	1.09
dystrophin myotonic-protein kinase	M87313	1.08
killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 5	AF217487	1.08
Consensus includes gb:AV781453 /FEA=EST /DB_XREF=gi:10919301 /DB_XREF=est:AV781453 /CLONE=M08BZA03 /UG=Hs.294014 ESTs	AV781453	1.08
egl-like module containing, mucin-like, hormone receptor-like sequence 1	NM_005430	1.08
whigless-type MMTV integration site family, member 1	NM_002472	1.08
myosin, heavy polypeptide 8, skeletal muscle, perinatal	BF593509	1.08
myosin binding protein C, slow type	AL567808	1.08
zinc finger protein 23 (K0X 16)	NM_033211	1.08
thymine-DNA glycosylase	NM_017911	1.08
hypothetical protein FLJ20635	NM_004796	1.08
neurexin 3	NM_006507	1.08
regenerating islet-derived 1 beta (pancreatic stone protein, pancreatic thread protein)	U06117	1.08
xanthine dehydrogenase	NM_018104	1.08
hypothetical protein FLJ10474	NM_001684	1.08
ATPase, Ca++ transporting, plasma membrane 4	NM_018118	1.08
hypothetical protein FLJ10508	NM_008643	1.08
serologically defined colon cancer antigen 3	BF692958	1.08
son of sevenless (Orsophilla) homolog 2	BF432532	1.08
TIA1 cytotoxic granule-associated RNA binding protein-like 1	NM_020127	1.08
tubulin 1	NM_014883	1.08
KIAA0914 gene product	NM_018593	1.08
hypothetical protein PRO0813	NM_024492	1.08
apolipoprotein (a) related gene C	NM_001340	1.08
cyclin, basic protein of sperm head cytoskeleton 2	AF070613	1.08
lankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase	NM_014840	1.08
KIAA0537 gene product	AW170549	1.08
hypothetical protein PRO2325	K03205	1.08
proline-rich protein Bat1 subfamily 2	AA142942	1.08
Consensus includes gb:AA142942 /FEA=EST /DB_XREF=gi:1712320 /DB_XREF=est:z43c04.s1 /CLONE=IMAGE:504678 /UG=Hs.241507 ribosomal protein S6	AA821844	1.08
DiGeorge syndrome gene A	U82317	1.08
arylsulfatase A	AB042825	1.08
RecQ protein-like 5	AU157008	1.08
proteasome (prosome, macropain) 26S subunit, non-ATPase, 5	AB011137	1.08
KIAA0565 gene product	NM_017490	1.08
ELKL motif kinase	NM_022058	1.08
solute carrier family 4, sodium bicarbonate cotransporter-like, member 10	U79302	1.08
p53-induced protein	BG403790	1.08
Consensus includes gb:BG403790 /FEA=EST /DB_XREF=gi:13287238 /DB_XREF=est:602419627F1 /CLONE=IMAGE:4526599 /UG=Hs.158154 ESTs	AL031848	1.08
cytosolic acyl coenzyme A thioester hydrolase	AI922519	1.08
rad8 GTPase activating protein (GAP and centrosome-associated)	NM_006880	1.08
transcription termination factor, mitochondrial	NM_015957	1.08
CGI-29 protein	AF261091	1.08
ATP-binding cassette, sub-family F (GCON20), member 2	AF112121	1.08
Consensus includes gb:AU146809 /FEA=EST /DB_XREF=gi:11008330 /DB_XREF=est:AU146809 /CLONE=HEMBB1001564 /UG=Hs.287476 Homo sapiens cDNA FLJ121	NM_018394	1.08
hypothetical protein FLJ11342	AL080169	1.08
DKFZP434C171 protein	AF082559	1.08
lankyrase, TRF1-interacting ankyrin-related ADP-ribose polymerase	AW969803	1.08
Consensus includes gb:AW969803 /FEA=EST /DB_XREF=gi:8159647 /DB_XREF=est:EST381881 /UG=Hs.105791 ESTs	L22454	1.08
nuclear respiratory factor 1	AF119868	1.08
retinol binding protein 4, plasma	NM_022785	1.08
hypothetical protein FLJ23588	NM_017789	1.08
hypothetical protein FLJ20392	NM_0159053	1.08
Consensus includes gb:AI590063 /FEA=EST /DB_XREF=gi:4599101 /DB_XREF=est:t75d05.x1 /CLONE=IMAGE:2224137 /UG=Hs.235160 Homo sapiens mRNA, cDNA	BE586776	1.08
similar to KIAA0010 gene product (H. sapiens)	NM_018650	1.08
MAP/microtubule affinity-regulating kinase 1	NM_018295	1.08
hypothetical protein FLJ11000	NM_0149710	1.08
Consensus includes gb:AL049710 /DEF=Human DNA sequence from clone 633H17 on chromosome 1p31.2-32.2. Contains a pseudogene similar to part of MTCO1 (Cyto	AW007532	1.08
Consensus includes gb:AW007532 /FEA=EST /DB_XREF=gi:5856310 /DB_XREF=est:ws52h07.x1 /CLONE=IMAGE:3500861 /UG=Hs.103391 Human insulin-like growth fr	NM_017874	1.08
hypothetical protein FLJ20550	NM_007036	1.08
endothelial cell-specific molecule 1		

carbonic anhydrase IV
 HMG-box containing protein 1
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 3
 KIAA0894 gene product
 growth hormone secretagogue receptor
 protein tyrosine phosphatase, non-receptor type substrate 1
 melanoma cell adhesion molecule
 Consensus includes gb:AL136527 /DEF=Human DNA sequence from clone RP11-215B13 on chromosome 13 Contains the AKAP11 (A kinase (PRKA) anchor protein 11);
 Consensus includes gb:AV652328 /FEA=EST /DB_XREF=gi:3873342 /DB_XREF=est:AV652328 /CLONE=GLCZC01 /UG=Hs.282488 ESTs
 calmodulin 1 (phosphorylase kinase, delta)
 hypothetical protein FLJ23403
 glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6-N-acetylglucosaminyltransferase)
 WW domain-containing oxidoreductase
 hypothetical protein FLJ11556
 Cluster Ind. N80935:207g06.a1 Homo sapiens cDNA, 3' end /clone=IMAGE:301402 /clone_end=3 /gb=N80935 /gi=1243636 /ug=Hs.22483 /len=527
 Consensus includes gb:AL138781 /DEF=Human DNA sequence from clone RP11-18H23 on chromosome 10. Contains the gene KIAA0204 (HSLX) for a protein kinase, thr
 corticotropin releasing hormone
 estrogen receptor 1
 major histocompatibility complex, class I, E
 geranylgeranyl diphosphate synthase 1
 lactase
 zinc finger protein 237
 lectin, galactoside-binding, soluble, 4 (galectin 4)
 Fc fragment of IgA, receptor for
 osteoglycin (osteoblast growth factor, mimetic)
 Consensus includes gb:AU145411 /FEA=EST /DB_XREF=gi:1106932 /DB_XREF=est:AU145411 /CLONE=HEMBA1004753 /UG=Hs.274546 Homo sapiens cDNA FLJ10
 KIAA0869 protein
 transcription factor 3 (E2A Immunoglobulin enhancer binding factors E12/E47)
 galactose-1-phosphate uridylyltransferase
 hypothetical protein FLJ20802
 hypothetical protein MGC2383
 Consensus includes gb:AK023690.1 /DEF=Homo sapiens cDNA FLJ13628 fts, clone PLACE1011054 /FEA=mRNA /DB_XREF=gi:10435692 /UG=Hs.306656 Homo sapier
 hypothetical protein MGC3035
 patched homolog 2 (Drosophila)
 small inducible cytokine A5 (RANTES)
 KIAA1040 protein
 hypothetical protein FLJ12572
 Consensus includes gb:AK024890.1 /DEF=Homo sapiens cDNA: FLJ21237 fts, clone COL01114 /FEA=mRNA /DB_XREF=gi:10437303 /UG=Hs.308720 Homo sapiens cD
 tumor necrosis factor receptor superfamily, member 18
 sterol O-acetyltransferase 2
 Fanconi anemia, complementation group F
 calcium channel, voltage-dependent, gamma subunit 1
 fibrinogen silencer binding protein
 Consensus includes gb:AK027246.1 /DEF=Homo sapiens cDNA: FLJ23593 fts, clone LNG14821 /FEA=mRNA /DB_XREF=gi:10440333 /UG=Hs.287749 Homo sapiens cD
 hypothetical protein FLJ11850
 pregnancy specific beta-1-glycoprotein 1
 ribosomal protein L13
 Consensus includes gb:AL133571.1 /DEF=Homo sapiens mRNA: cDNA DKFZp434F1135 (from clone DKFZp434F1135) /FEA=mRNA /DB_XREF=gi:5599148 /UG=Hs.11
 acetylcholinesterase (YT blood group)
 hypothetical protein MGC2815
 T cell receptor gamma locus
 phosphatidylinositol transfer protein
 T cell receptor gamma constant 2
 cyclic AMP phosphoprotein, 19 kD
 hypothetical protein FLJ10241
 Huntingtin interacting protein E
 f-box and leucine-rich repeat protein 11
 telomerase-associated protein 1
 paired box gene 8
 canine palmitoyltransferase II
 thrombospondin 1
 interleukin 1, alpha
 GM2 ganglioside activator protein

heme binding protein 1
zinc finger protein, subfamily 1A, 5 (Pegasus)
Consensus includes gb:BC005286.1 /DEF=Homo sapiens, clone IMAGE3935812, mRNA, partial cds. /FEA=mRNA /PROD=Unknown (protein for IMAGE3935812) /DB_Xt BC005286
chromosome 21 open reading frame 18
hypothetical protein FLJ22402
gb:AF276653.1 /DEF=Homo sapiens MAP1 light chain 3-like protein 2 mRNA, complete cds. /FEA=mRNA /PROD=MAP1 light chain 3-like protein 2 /DB_XREF=gi:1362577 AF276653
Consensus includes gb:AL359599.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547C126 (from clone DKFZp547C126). /FEA=mRNA /DB_XREF=gi:8655666 /UG=Hs.2838 AL359599
gamma-aminobutyric acid (GABA) B receptor, 1
Consensus includes gb:AL390143.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547N074 (from clone DKFZp547N074). /FEA=mRNA /DB_XREF=gi:9368876 /UG=Hs.2842 AL390143
tripartite motif-containing 15
histone deacetylase 6
cell division cycle 25A
Consensus includes gb:AL137285.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D2416 (from clone DKFZp434D2416). /FEA=mRNA /DB_XREF=gi:6807740 /UG=Hs.30 AL137285
pregnancy-associated plasma protein A
occludin
protein phosphatase 5, catalytic subunit
hypothetical protein FLJ10040
period homolog 2 (Drosophila)
basic transcription factor 3
p53-regulated DDA3
Consensus includes gb:AL049646 /DEF=Human DNA sequence from clone RP4-568F9 on chromosome 20 Contains the ZNF133 (zinc finger protein 133 (clone pHZ-13)) /AL049646
DKFZP547G0910 protein
phosphatase and tetraol homolog (mutated in multiple advanced cancers 1)
zinc finger protein, Y-linked
Kallmann syndrome 1 sequence
glutamate-aminotransferase (glutamine synthase)
keratin 18
aspartate beta-hydroxylase
thrombosane A2 receptor
RAS, member RAS oncogene family
guanosine monophosphate reductase
plectin 1, intermediate filament binding protein, 500kD
transferrin
chitinase 1 (chitinohydrolase)
hypothetical protein MGC4365
GRB2-associated binding protein 2
protein kinase, AMP-activated, alpha 1 catalytic subunit
PHD finger protein 1
gb:AF130050.1 /DEF=Homo sapiens clone FLB3442 PRO0872 mRNA, complete cds. /FEA=mRNA /PROD=PRO0872 /DB_XREF=gi:11493406 /UG=Hs.172154 Homo sapiens
acrosin
phosphatidylinositol-4-phosphate 5-kinase, type II, beta
KIAA1116 protein
KIAA0346 protein
testis transcript Y 9
KIAA0449 protein
Consensus includes gb:AU146999 /FEA=EST /DB_XREF=gi:11008520 /DB_XREF=est:AU146999 /CLONE=HEMBB1002092 /UG=Hs.296688 Homo sapiens cDNA FLJ121
Consensus includes gb:AL577024 /FEA=EST /DB_XREF=gi:12939748 /DB_XREF=est:AL577024 /CLONE=CS001082YN19 (3 prime) /UG=Hs.7921 Homo sapiens mRNA; AL577024
DKFZP34K114 protein
carcinoembryonic antigen-related cell adhesion molecule 6 (non-specific cross reacting antigen)
keratin, hair, basic, 5
p53 regulated PA28 nuclear protein
KIAA1416 protein
ATP-binding cassette, sub-family A (ABC1), member 8
similar to hypothetical protein MNCb-2386
tumor endothelial marker 8
statin 1/ontoprotein 18
splicing factor proline/glutamine rich (polypyrimidine tract binding protein associated)
hypothetical protein FLJ20371
homologous to yeast nitrogen permease (candidate tumor suppressor)
calcium channel, voltage-dependent, alpha 1G subunit
hypothetical protein MGC13033
gb:AF212224.1 /DEF=Homo sapiens CLK4 mRNA, complete cds. /FEA=mRNA /PROD=CLK4 /DB_XREF=gi:9437514 /UG=Hs.295231 Homo sapiens CLK4 mRNA, compk AF212224
tropomyosin 1 (alpha)

stromal cell protein
 Consensus includes gb:8E613017 /FEA=EST /DB_XREF=gi:10245251 /DB_XREF=est:RC2-AN0062-140800-017-g03 /UG=Hs.124629 ESTs
 hypothetical protein FLJ20514
 Consensus includes gb:AK025194.1 /DEF=Homo sapiens cDNA: FLJ21541 flt. clone COL08166. /FEA=mRNA /DB_XREF=gi:10437857 /UG=Hs.308784 Homo sapiens cDNA: AK025194
 retinoic acid repressible protein
 Consensus includes gb:AV711904 /FEA=EST /DB_XREF=gi:10731210 /DB_XREF=est:AV711904 /CLONE=DCAAIE08 /UG=Hs.277431 Homo sapiens cDNA: FLJ23356 flt. AV711904
 Protein P3
 gb:BC004241.1 /DEF=Homo sapiens. Similar to laminin, alpha 4, clone MGC:10389, mRNA, complete cds. /FEA=mRNA /PROD=Similar to laminin, alpha 4 /DB_XREF=gi: BC004241
 KIAA0222 gene product
 AL110158
 U04845
 collagen, type IV, alpha 6
 U04845
 sarcoplasmic
 hypothetical protein FLJ21820
 quinine oxidoreductase homolog
 PAS-serine/threonine kinase
 trefoil factor 2 (spasmolytic protein 1)
 KIAA0878 protein
 ELL-RELATED RNA POLYMERASE II, ELONGATION FACTOR
 host cell factor homolog
 BCL2-interacting killer (apoptosis-inducing)
 ISL1 transcription factor, LIMhomeodomain, (Islet-1)
 signal-regulatory protein beta 1
 Consensus includes gb:AF217536.1 /DEF=Homo sapiens truncated mevalonate kinase mRNA, partial cds, alternatively spliced. /FEA=mRNA /PROD=truncated mevalonate
 GS3955 protein
 myotubularin related protein 2
 syndecan 4 (amphiglycan, ryudocan)
 arginase, type II
 neurotrophic tyrosine kinase, receptor, type 2
 actin cross-linking factor
 PRO1575 protein
 Consensus includes gb:AW451236 /FEA=EST /DB_XREF=gi:6992012 /DB_XREF=est:UI-H-B13-ah-a-07-0-UI.s1 /CLONE=IMAGE:2736661 /UG=Hs.15535 Homo sapiens
 hypothetical protein FKSG44
 protocadherin gamma subfamily A, 3
 alkaline phosphatase, placental (Regan isozyme)
 phosphatidylinositol glycan, class F
 solute carrier family 19 (folate transporter), member 1
 UHS KerB
 v-Ki-ras2 Kirsten rat sarcoma 2 viral oncogene homolog
 hypothetical protein FLJ20059
 fragile X mental retardation, autosomal homolog 2
 Consensus includes gb:AW972855 /FEA=EST /DB_XREF=gi:8162701 /DB_XREF=est:EST384950 /UG=Hs.292853 ESTs
 phosphate cytidylyltransferase 1, choline, alpha isoform
 cytochrome P450, subfamily IIA (phenobarbital-inducible), polypeptide 13
 c-myc promoter-binding protein
 TAR (HIV) RNA binding protein 1
 hypothetical protein FLJ23407
 hypothetical protein FLJ13687
 thrombospondin 1
 forkhead box M1
 fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson
 Consensus includes gb:AV194543 /FEA=EST /DB_XREF=gi:6473381 /DB_XREF=est:b27c02.s1 /CLONE=IMAGE:2577506 /UG=Hs.234573 Homo sapiens mRNA for TL AW194543
 hypothetical protein FLJ22823
 zinc finger protein 35 (clone HF.10)
 KIAA0663 gene product
 KIAA0752 protein
 Cluster Ind. N36928; yy3806.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:273538 /clone_end=3 /gb=N36928 /gi=1158070 /ug=Hs.33540 /len=562
 KIAA0356 gene product
 Consensus includes gb:AF007147.1 /DEF=Homo sapiens clone 23712 mRNA sequence. /FEA=mRNA /DB_XREF=gi:2852625 /UG=Hs.113166 Homo sapiens clone 23712 AF007147
 HNK-1 sulfotransferase
 cholinergic receptor, nicotinic, alpha polypeptide 10
 myotubular myopathy 1
 polymerase I and transcript release factor
 interleukin 11

gb:AL136548.1 /DEF=Homo sapiens mRNA; cDNA DKFZp761G18121 (from clone DKFZp761G18121); complete cds. /FEA=mRNA /GEN=DKFZp761G18121 /PROD=hyp
 pancreatic elastase 1B
 1.08 NM_015849
 hypothetical protein FLJ20070
 1.08 NM_017652
 Consensus includes gb:U00773.1 /DEF=Human EST clone 42944 mariner transposon Hsmar1 sequence /FEA=mRNA /DB_XREF=gi:2231376 /UG=Hs.121580 Human ES
 U00773
 1.08 BG111661
 gqgi autanigen, gqgin subfamily a. 1
 1.08 NM_008080
 gamma-aminobutyric acid (GABA) A receptor, alpha 3
 1.08 NM_014237
 a disintegrin and metalloproteinase domain 18
 1.08 AWT48405
 neurexin 1
 1.08 NM_030808
 LIS1-interacting protein NUDEL; endonucleopeptidase A
 1.08 NM_007350
 pleckstrin homology-like domain, family A, member 1
 1.08 BC001090
 KIAA1668 protein
 1.08 AB014607
 KIAA0707 protein
 1.08 NM_016265
 G1OT-3 for gonadotropin inducible transcription repressor-3
 1.08 AG02100
 dedual protein induced by progesterone
 1.08 AF068266
 core-binding factor, runt domain, alpha subunit 2; translocated to, 2
 1.08 NM_006606
 retinoblastoma binding protein 9
 1.08 AW007137
 hypothetical protein FLJ10342
 1.08 NM_003186
 transgelin
 1.08 NM_004726
 RALBP1 associated Eps domain containing 2
 1.08 NM_024300
 hypothetical protein MGC2217
 1.08 NM_001297
 cyclic nucleotide gated channel beta 1
 1.08 NM_002114
 human immunodeficiency virus type 1 enhancer binding protein 1
 1.08 AA911235
 macrophage stimulating, pseudogene 9
 1.08 NM_001255
 CDC20 cell division cycle 20 homolog (S. cerevisiae)
 1.08 AJ133788
 LIM domain binding 3
 1.08 NM_025257
 NG22 protein
 1.08 AA723370
 CGI-105 protein
 1.08 A073549
 estrogen receptor 1
 1.08 Z70519
 tumor necrosis factor receptor superfamily, member 6
 1.08 NM_024759
 hypothetical protein FLJ13955
 1.08 NM_0110164
 Consensus includes gb:AL110164.1 /DEF=Homo sapiens mRNA; cDNA DKFZp58610324 (from clone DKFZp58610324). /FEA=mRNA /DB_XREF=gi:5817069 /UG=Hs.1937
 AL110164
 1.08 NM_018614
 hypothetical protein PRO2012
 1.08 AA422049
 hypothetical protein DKFZp547M138 similar to widely-interspaced zinc finger motifs
 1.08 AK022326
 Consensus includes gb:AK022326.1 /DEF=Homo sapiens cDNA FLJ12264 fls. clone MAMMA1001600. /FEA=mRNA /DB_XREF=gi:10433699 /UG=Hs.288481 Homo sapie
 trophin
 1.08 AF349719
 dystrobrevin, alpha
 1.08 NM_001390
 ubiquitin specific protease 24
 1.08 BF444943
 cholecystokinin A receptor
 1.08 L13605
 tetraspan 1
 1.08 AF133425
 KIAA1464 protein
 1.08 AL041852
 bubby like protein 1
 1.08 NM_003322
 serine (or cysteine) proteinase inhibitor, clade D (heparin cofactor), member 1
 1.08 NM_000165
 Consensus includes gb:AL139112 /DEF=Human DNA sequence from clone GS1-103B18 on chromosome Xq27.1-27.3 Contains ESTs, STSs and GSSs. Contains a novel
 mucin 8, tracheobronchial
 1.08 AL139112
 HGC8.1.1 protein
 1.08 U14383
 bone gamma-carboxyglutamate (gla) protein (osteocalcin)
 1.08 AB016900
 exotoxins (multiple)-like 3
 1.07 NM_000711
 trinuclidean repeat containing 15
 1.07 AF001690
 hypothetical protein FLJ21497
 1.07 AL045800
 thyroid hormone receptor-associated protein, 95-kD subunit
 1.07 NM_025102
 CATX-8 protein
 1.07 AF106934
 surfactant, pulmonary-associated protein C
 1.07 NM_020387
 thyroid hormone receptor coactivating protein
 1.07 AA633841
 nuclear receptor subfamily 2, group C, member 1
 1.07 AL136823
 hypothetical protein MGC3265
 1.07 NM_003297
 lamin A/C
 1.07 NM_024028
 Consensus includes gb:AK026894.1 /DEF=Homo sapiens cDNA: FLJ23041 fls. clone LNG02288. /FEA=mRNA /DB_XREF=gi:10439604 /UG=Hs.306865 Homo sapiens cC
 AK026894
 1.07 NM_001488
 glutamate-cysteine ligase, catalytic subunit
 1.07 BC000076
 cyclin D1 (PRAD1; parathyroid adenomatosis 1)
 1.07 AK021487
 Consensus includes gb:AK021487.1 /DEF=Homo sapiens cDNA FLJ11425 fls. clone HEMBA1001051. /FEA=mRNA /DB_XREF=gi:10432680 /UG=Hs.306597 Homo sapie
 Ketch motif containing protein
 1.07 BC005253
 KIAA0876 protein
 1.07 AK026040
 Notch homolog 4 (Drosophila)
 1.07 A1743713

casepase 9, apoptosis-related cysteine protease	U60521	1.07
tripartite motif-containing 15	U34249	1.07
transforming growth factor, beta receptor I (activin A receptor type II-like kinase, 53KD)	NM_004612	1.07
decdinase, dcdomyonine, type III	NM_001382	1.07
succinate-CoA ligase, GDP-forming, beta subunit	AV650470	1.07
ATP-binding cassette, sub-family D (ALD), member 4	NM_005050	1.07
serine/threonine kinase 17a (apoptosis-inducing)	AW183478	1.07
hypothetical protein FLJ14360	BG325646	1.07
myosin, light polypeptide 5, regulatory	L03785	1.07
contactin associated protein-like 2	AU144598	1.07
PAB23, member PAS oncogene family	NM_016277	1.07
estrogen regulated gene 1	NM_022034	1.07
cadherin 6, type 2, K-cadherin (fetal kidney)	AU151463	1.07
catenin, beta interacting protein 1	NM_020248	1.07
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump) membrane sector associated protein M8-9	AF248968	1.07
neuronal pentraxin receptor	AL162057	1.07
XO4603 B subunit thrc, thrb genes corresponding to nucleolides 248-2228 of XO4603 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)	XO4603	1.07
Consensus includes gb:BF223370 /FEA=EST /DB_XREF=gi:11130547 /DB_XREF=est:q87603.x1 /CLONE=IMAGE:3705316 /UG=Hs.90449 Human clone 23908 mRNA	BF223370	1.07
phosphatidylserine-specific phospholipase A1 alpha	NM_015900	1.07
excision repair cross-complementing rodent repair deficiency, complementation group 6	NM_000124	1.07
insulin-like 4 (placenta)	NM_002195	1.07
Consensus includes gb:U79249.1 /DEF=Human clone 23839 mRNA sequence. /FEA=mRNA /DB_XREF=gi:1710196 /UG=Hs.76362 Human clone 23839 mRNA sequence	U79249	1.07
ribosomal protein S6 kinase, 80KD, polypeptide 6	AF184965	1.07
chordin-like	AL049176	1.07
protocadherin gamma subfamily C, 3	NM_002588	1.07
transferrin receptor (p90, CD71)	BC001188	1.07
tumor protein p53 binding protein, 1	NM_005657	1.07
metallothionein 3 (growth inhibitory factor (neurotrophic))	NM_005954	1.07
interleukin 7	NM_000880	1.07
pyrimidinergic receptor P2Y, G-protein coupled, 4	NM_002585	1.07
hypothetical protein	NM_016101	1.07
lymphocyte antigen 8 complex, locus D	NM_003695	1.07
vav 3 oncogene	AF118887	1.07
leukocyte-associated Ig-like receptor 2	NM_002288	1.07
KIA0381 protein	BF513244	1.07
KIA0197 protein	D83781	1.07
hypothetical protein MGC2963	NM_031298	1.07
KIA11718 protein	BE217882	1.07
serine/threonine kinase 38 like	AW779556	1.07
B-cell CLL/lymphoma 7A	BC002829	1.07
Integrin, beta 4	AA808063	1.07
exostoses (multiple)-like 2	AF000416	1.07
ubiquitin specific protease 21	NM_012475	1.07
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila), translocated to, 3	AV756536	1.07
kinase-like 7	NM_020242	1.07
Consensus includes gb:AU145711 /FEA=EST /DB_XREF=gi:11007232 /DB_XREF=est:AU145711 /CLONE=HEMBA1005588 /UG=Hs.301006 Homo sapiens cDNA FLJ11	AU145711	1.07
melanoma antigen, family A, 12	BC003408	1.07
plihn, desmosome associated protein	BF508848	1.07
transitional epithelia response protein	NM_013319	1.07
PTH-responsive osteosarcoma B1 protein	U87408	1.07
Interleukin 6 signal transducer (gp130, oncostatin M receptor)	BE856546	1.07
DKF ZP434N014 protein	AA813332	1.07
single-strand selective monofunctional uracil DNA glycosylase	NM_014311	1.07
C-terminal binding protein 1	BF337195	1.07
myosin, heavy polypeptide 2, skeletal muscle, adult	NM_017534	1.07
STO protein	NM_014070	1.07
casepase 10, apoptosis-related cysteine protease	U88214	1.07
Consensus includes gb:AL157375 /DEF=Human DNA sequence from clone RP1-133P16 on chromosome 6 Contains the ARF1 (ADP-ribosylation factor 1) gene, ESTs, ST. AL157375	AL157375	1.07
hypothetical protein FLJ21472	NM_024837	1.07
Consensus includes gb:AK0222363.1 /DEF=Homo sapiens cDNA FLJ12301 fls, clone MAMMA1001858. /FEA=mRNA /DB_XREF=gi:10433745 /UG=Hs.287511 Homo sapie	AK022363	1.07
dihydropyrate dehydrogenase	BE789211	1.07
hypothetical protein	NM_016399	1.07
KIAA0707 protein	AK023537	1.07

beta-1,4 mannosyltransferase
Consensus includes gb:AF103295.1 /DEF=Homo sapiens clone N87 immunoglobulin heavy chain variable region mRNA, partial cds. /FEA=mRNA /PROD=immunoglobulin
NM_004148 1.07

ninjurin 1
AE binding protein 1
KIAA0962 protein
complement component 3
tumor necrosis factor (ligand) superfamily, member 6
hypothetical protein FLJ12377
phosphatidylinositol glycan, class L
G protein-coupled receptor 25
NM_000064 1.07

hypothetical protein FLJ11743
Consensus includes gb:AA875427 /FEA=EST /DB_XREF=gi:3151219 /DB_XREF=estco28g02.s1 /CLONE=IMAGE:1587698 /UG=Hs.128434 Homo sapiens ELISC-1 mRN
NM_024527 1.07

solute carrier family 2 (facilitated glucose transporter), member 3
sphingosine-1-phosphate lyase 1
small inducible cytokine subfamily B (Cys-X-Cys), member 10
CGI-44 protein; sulfide dehydrogenase like (yeast)
catenin (cadherin-associated protein), alpha-like 1
nuclear receptor subfamily 5, group A, member 2
golgi-specific brefeldin A resistance factor 1
caspase 10, apoptosis-related cysteine protease
KIAA0202 protein
NM_004163 1.07

p10-binding protein
myosin, heavy polypeptide 11, smooth muscle
protocadherin gamma subfamily A, 9
protein kinase C binding protein 1
carbohydrate (keratan sulfate Gal-6) sulfotransferase 1
cholinergic receptor, nicotinic, delta polypeptide
superficial viralidic activity 2-like (S. cerevisiae)
deleted in lung and esophageal cancer 1
breakpoint cluster region
vasoactive intestinal peptide receptor 2
enhancer of rudimentary homolog (Drosophila)
phosphoinositide-specific phospholipase C-beta 1
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson
AB030078 1.07

Pleuroctosomal GTP-binding protein-like
seven transmembrane protein TM7SF3
uncharacterized hematopoietic stem/progenitor cells protein MDS028
nucleosome assembly protein 1-like 1
glutamate receptor, ionotropic, N-methyl D-aspartate 1
spinocerebellar ataxia 8
terranectin (plasminogen binding protein)
disrupted in bipolar disorder 1
deleted in cancer 1; RNA helicase HDB/DICE1
KIAA0274 gene product
UDP glucosyltransferase 2 family, polypeptide A1
ring finger protein 15
G protein-coupled receptor kinase 2-like (Drosophila)
KIAA0210 gene product
NM_004537 1.07

hypothetical protein FLJ11827
Consensus includes gb:AP000693 /DEF=Homo sapiens genomic DNA, chromosome 21q22.2, PCR fragment from BAC clone:XB739C11, CBR1-HLCS region /FEA=mRNA /P
A0920693 1.07

KIAA1117 protein
TATA box binding protein (TBP)-associated factor, RNA polymerase I, A, 48kD
Golgi apparatus protein 1
ecotropic viral integration site 1
PMAS2 postmeiotic segregation increased 2 (S. cerevisiae)
keratin, hair, acidic, 1
calumenin
retinol X receptor, beta
CGI-48 protein
mannose-binding lectin (protein C) 2, soluble (opsonic defect)
polymerase (RNA mitochondrial [DNA directed])
BF057617 1.07

synaptrophin, alpha 1(dystrophin-associated protein A1, 59kD, acidic component)
 epidermal growth factor receptor pathway substrate 8
 hypothetical protein FLJ12078
 thoredoxin domain-containing
 integrin, beta 3 (platelet glycoprotein IIb, antigen CD61)
 hypothetical protein FLJ21308
 Consensus includes gb:AL390137.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434D179 (from clone DKFZp434D179). /FEA=mRNA /DB_XREF=gi:9368847 /UG=Hs.3065
 belaine-homocysteine methyltransferase 2
 purinergic receptor P2X, ligand-gated ion channel, 1
 LIM homeobox protein 8
 homeodomain interacting protein kinase 3
 cardioembryonic antigen-related cell adhesion molecule 7
 nudaportin 88KD
 contactin 5
 p53-induced protein PIGPC1
 activin A receptor type II-like 1
 UDP-glucose ceramide glucosyltransferase-like 2
 KIAA1503 protein
 aquaporin 5
 claudin 16
 PHD finger protein 1
 Consensus includes gb:AL137713.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434I0523 (from clone DKFZp434I0523). /FEA=mRNA /DB_XREF=gi:6808127 /UG=Hs.3064 /AL137713
 amyloid beta (A4) precursor protein-binding, family A, member 2 (X11-like)
 gb:NM_003902.1 /DEF=Homo sapiens far upstream element (FUSE) binding protein 1 (FUBP1), mRNA. /FEA=mRNA /GEN=FUBP1 /PROD=far upstream element-binding
 BC000585
 solute carrier family 21 (organic anion transporter), member 11
 R3H domain (binds single-stranded nucleic acids) containing
 plastin 3 (T isoform)
 apolipoprotein L, 3
 adrenal gland protein AD-004
 KIAA0096 protein
 A1Pase, Cs++ transporting, plasma membrane 3
 activating transcription factor 5
 PTH-responsive osteosarcoma B1 protein
 popeye protein 3
 novel RGD-containing protein
 hypothetical protein FLJ13852
 RAB2, member RAS oncogene family
 Consensus includes gb:A1769688 /FEA=EST /DB_XREF=gi:5236197 /DB_XREF=est:W2511.x1 /CLONE=IMAGE:2403885 /UG=Hs.180737 Hmo sapiens clone 23664 an
 A1769688
 hypothetical protein FLJ11585
 KIAA0232 gene product
 DnaJ (Hsp40) homolog, subfamily C, member 8
 BN51 (BHK21) temperature sensitivity complementing
 solute carrier family 31 (copper transporters), member 1
 reversion-inducing-cysteine-rich protein with kazal motifs
 KIAA0805 protein
 M24537B subtilis pheA genes corresponding to nucleotides 2017-3334 of M24537 (-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 prime respectively)
 frizzled homolog 8 (Drosophila)
 Consensus includes gb:A1471375 /FEA=EST /DB_XREF=gi:4333465 /DB_XREF=est:tm10c12.x1 /CLONE=IMAGE:2156182 /UG=Hs.80206 glucose-6-phosphate dehydrog
 A471375
 thrombospondin 3
 short stature homeobox 2
 mitochondrial ribosomal protein S26
 Consensus includes gb:AL137378.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434K1128 (from clone DKFZp434K1128). /FEA=mRNA /DB_XREF=gi:6807908 /UG=Hs.30
 AL137378
 CD38 antigen (collagen type I receptor, thrombospondin receptor)-like 1
 splicing factor 3b, subunit 4, 49KD
 putative translation initiation factor
 villin-like
 S100 calcium binding protein A7 (psoriasin 1)
 procollagen-lysine, 2-oxoglutarate 5-dioxygenase 3
 hypothetical protein FLJ11078
 apolipoprotein E
 cyclin T2
 tyrosine hydroxylase
 UDP-N-acetyl-alpha-D-galactosamine (N-acetylneuraminy)l-galactosylglucosylceramide N-acetylgalactosaminyltransferase (GalNAc-T)
 ALD50358

Rho guanine exchange factor (GEF) 11
 Consensus includes gb:AK024688.1 /DEF=Homo sapiens cDNA: FLJ12151 fs, clone COL00526, /FEA=mrRNA /DB_XREF=gi:10437278 /UG=Hs.306711 Homo sapiens cDNA: AK024688
 carboxypeptidase E
 Consensus includes gb:AK022303.1 /DEF=Homo sapiens cDNA: FLJ12241 fs, clone MAMMA1001274, /FEA=mrRNA /DB_XREF=gi:10433670 /UG=Hs.287503 Homo sapiens cDNA: AK022303
 erythroid differentiation and denudation factor 1
 DNFXP34A0131 protein
 WNT1 inducible signaling pathway protein 1
 deleted in malignant brain tumors 1
 isochlorate dehydrogenase 3 (NAD+) alpha
 GAS2-related on chromosome 22
 adenylyate cyclase 2 (brain)
 KIAA1464 protein
 caspase recruitment domain protein 14
 hypothetical protein similar to preferentially expressed antigen of melanoma
 nuclear receptor co-repressor 1
 ubiquitously transcribed tetratricopeptide repeat gene, Y chromosome
 Consensus includes gb:AU148184 /FEA=EST /DB_XREF=gi:11009705 /DB_XREF=est:AU148184 /CLONE=MAMMA1002820 /UG=Hs.287529 Homo sapiens cDNA: FLJ12 AU148184
 CS box-containing WD protein
 RAN binding protein 2-like 1
 H1 histone family, member 3
 Consensus includes gb:AU073983 /FEA=EST /DB_XREF=gi:3400627 /DB_XREF=est:oy6604.x1 /CLONE=IMAGE:1670790 /UG=Hs.298553 Homo sapiens, clone IMAGE: AU073983
 fatty acid desaturase 1
 cAMP responsive element modulator
 syntactin binding protein 3
 nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, beta
 PTH-responsive osteosarcoma B1 protein
 paired basic amino acid cleaving enzyme (furin, membrane associated receptor protein)
 hypothetical protein FLJ10955
 dopamine receptor D5
 pyruvate dehydrogenase kinase, isoenzyme 4
 apolipoprotein A-II
 Consensus includes gb:AL049244.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564C163 (from clone DKFZp564C163), /FEA=mrRNA /DB_XREF=gi:4499979 /UG=Hs.3020 AL049244
 Consensus includes gb:AU157200 /FEA=EST /DB_XREF=gi:11018721 /DB_XREF=est:AU157200 /CLONE=PLACE1006617 /UG=Hs.301390 Homo sapiens cDNA: FLJ135 S73854
 neurofibromin 2 (bilateral acoustic neuroma)
 neural polyrimidine tract binding protein
 galactose-4-epimerase, UDP-
 solute carrier family 7 (cationic amino acid transporter, y+ system), member 1
 collagen, type IV, alpha 6
 killer cell lectin-like receptor subfamily C, member 2
 tumor necrosis factor receptor superfamily, member 9
 hypothetical protein FLJ10378
 adrenergic, beta, receptor kinase 1
 hypothetical protein FLJ10252
 cystatin SA
 solute carrier family 6 (neurotransmitter transporter, dopamine), member 3
 KIAA1077 protein
 hypothetical protein FLJ10339
 agrin
 nucleoside peptide receptor C/guanylate cyclase C (atrionatriuretic peptide receptor C)
 cytokine receptor-like factor 2
 hypothetical protein FLJ20531
 Consensus includes gb:T53900 /FEA=EST /DB_XREF=gi:655761 /DB_XREF=est:y83e08.r1 /CLONE=IMAGE:77798 /UG=Hs.172035 hypothetical protein similar to mouse: T53900
 Consensus includes gb:AI382123 /FEA=EST /DB_XREF=gi:4194904 /DB_XREF=est:b30e09.x1 /CLONE=IMAGE:2087416 /UG=Hs.182919 ESTs, Highly similar to A5925: AI382123
 Rho guanine nucleotide exchange factor (GEF) 10
 zinc finger protein 225
 hypothetical protein MGC9084
 cholinergic receptor, nicotinic, alpha polypeptide 3
 chromodomain helicase DNA binding protein 1
 netrin G1
 NDRG family, member 4
 5-methyltetrahydrofolate-homocysteine methyltransferase reductase
 immature colon carcinoma transcript 1
 retinol dehydrogenase 5 (11-cis and 9-cis)

chromosome 1 open reading frame 16
 G protein-coupled receptor 51
 Consensus includes gb:AU147084 /FEA=EST /DB_XREF=est:AU147084 /CLONE=HEMBB1002371 /UG=Hs.287482 Homo sapiens cDNA FLJ121
 cameline palmitoyltransferase II
 protein tyrosine phosphatase, non-receptor type 21
 Ca2+-dependent activator protein for secretion
 cd.1A
 hemoglobin, theta 1
 serine (or cysteine) proteinase inhibitor, clade G (C1 inhibitor), member 1
 involucrin
 ADP-ribosyltransferase 1
 similar to RhoGAP
 DKFZP434C212 protein
 growth arrest-specific 1
 CGI-14 protein
 Consensus includes gb:AL359563.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434D0328 (from clone DKFZp434D0328) /FEA=mRNA /DB_XREF=gi:8655619 /UG=Hs.30
 immunoglobulin lambda joining 3
 suppressor of Ty 4 homolog 1 (S. cerevisiae)
 protocadherin beta 17 pseudogene
 Rhesus blood group, CcEe antigens
 Consensus includes gb:AL389866 /DEF=Human DNA sequence from clone RP1-237C24 on chromosome 20 Contains the 3 end of the SLC23A1 gene encoding solute car
 heparan sulfate (glucosamine) 3-O-sulfotransferase 2
 ADP-ribosylation factor-like 7
 KIAA0848 protein
 RAB31, member RAS oncogene family
 ring finger protein 11
 insulin-like growth factor binding protein 1
 insulin-like growth factor binding protein 1
 T cell receptor alpha locus
 RAB5C, member RAS oncogene family
 cyokeratin type II
 SRY (sex determining region Y)-box 9 (campomic dysplasia, autosomal sex-reversal)
 solute carrier family 28 (sodium-coupled nucleoside transporter), member 2
 hypothetical protein FLJ10143
 TANK-binding kinase 1
 KIAA0141 gene product
 like mouse brain protein E46
 putative G-protein coupled receptor
 potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2
 Consensus includes gb:AU144324 /FEA=EST /DB_XREF=gi:11005845 /DB_XREF=est:AU144324 /CLONE=HEMBA1001570 /UG=Hs.306599 Homo sapiens cDNA FLJ11
 transcription factor EC
 matrix metalloproteinase 19
 Cluster incl. A1W033081.wrQ3d04.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-2480455 /clone_end=3 /gb=AW003091 /gi=5849929 /ug=Hs.6217 /len=613
 EGF-like domain, multiple 5
 hypothetical gene CQ018
 retinol binding protein 1, cellular
 Consensus includes gb:AU153979 /FEA=EST /DB_XREF=gi:1015500 /DB_XREF=est:AU153979 /CLONE=NT2RP3004470 /UG=Hs.293928 Homo sapiens cDNA FLJ142
 zinc finger protein 235
 thyroid hormone receptor, beta (erythroblastic leukemia viral (v-erb-a) oncogene homolog 2, avian)
 KIAA0649 gene product
 Mouse Mammary Tumor Virus Receptor homolog
 deoxyribonuclease I-like 2
 KIAA0383 protein
 KIAA0472 protein
 heterogeneous nuclear ribonucleoprotein H2 (H')
 heparan sulfate 2-O-sulfotransferase
 collagen, type V, alpha 1
 transmembrane 4 superfamily member (tetraspan NET-7)
 cadherin 4, type 1, R-cadherin (relial)
 adrenergic, alpha-1D, receptor
 syndecan 3 (N-syndecan)
 KIAA0379 protein
 KIAA0335 gene product
 islet amyloid polypeptide

coagulation factor II (thrombin) receptor-like 3	NM_003950	1.06
collagen, type IV, alpha 3 (Goodpasture antigen)	M81379	1.06
KIAA0500 protein	AW028110	1.06
KIAA0523 protein	AL13046	1.06
KIAA0328 protein	AW003635	1.06
hypothetical protein PRO1188	NM_018621	1.06
Consensus includes gb:AW026481 /FEA=EST /DB_XREF=gi:5880011 /DB_XREF=est:ww14a04.x1 /CLONE=IMAGE:2529486 /UG=Hs.48396 ESTs, Moderately similar to	AW026481	1.06
regulator of G-protein signalling 4	NM_005613	1.06
SCN Circadian Oscillatory Protein (SCOP)	AB011178	1.06
short stature homeobox	NM_000451	1.06
aquaporin 3	AB001325	1.06
tripartite motif-containing 15	U34249	1.06
hypothetical protein PRO1598	NM_018503	1.06
caspase 8, apoptosis-related cysteine protease	NM_001228	1.06
gliome-amplified sequence-41	NM_006530	1.06
chloride channel 3	AA802571	1.06
chromosome 6 open reading frame 35	NM_018452	1.06
G protein-coupled receptor 24	NM_005297	1.06
hypothetical protein FLJ20116	AA469071	1.06
gb:AF130053.1 /DEF=Homo sapiens clone FLB4228 PRO1095 mRNA, complete cds, /FEA=mRNA /PROD=PRO1095 /DB_XREF=gi:11493412 /UG=Hs.302145 Homo sap	AF130053	1.06
teratocapsula repeat domain 3	NM_003316	1.06
tumor protein p53 binding protein, 2	NM_005426	1.06
caspace 7, apoptosis-related cysteine protease	NM_001227	1.06
jerky homolog (mouse)	AF072486	1.06
homeo box D3	NM_008888	1.06
KIAA0229 protein	AI990523	1.06
potassium voltage-gated channel, KOT-like subfamily, member 1	NM_000218	1.06
Consensus includes gb:N31716 /FEA=EST /DB_XREF=gi:1152115 /DB_XREF=est:yl5c12.s1 /CLONE=IMAGE:271318 /UG=Hs.31754 ESTs	N31716	1.06
Consensus includes gb:AU118026 /FEA=EST /DB_XREF=gi:109330043 /DB_XREF=est:AU118026 CLONE=HEMBA1002729 /UG=Hs.289068 Homo sapiens cDNA, FLJ11	AU118026	1.06
Consensus includes gb:AW300309 /FEA=EST /DB_XREF=gi:5709909 /DB_XREF=est:xs59608.x1 /CLONE=IMAGE:2773958 /UG=Hs.27453 ESTs, Moderately similar to	AW300309	1.06
carbonic anhydrase IX	NM_001216	1.06
potassium inwardly-rectifying channel, subfamily J, member 5	D50134	1.06
hypothetical protein MGC2869	BC005856	1.06
protein tyrosine phosphatase, non-receptor type 4 (megakaryocyte)	NM_002830	1.06
rhodopsin (opsin 2, rod pigment) (retinitis pigmentosa 4, autosomal dominant)	AA058638	1.06
GS15	NM_016526	1.06
Consensus includes gb:R25692 /FEA=EST /DB_XREF=gi:761827 /DB_XREF=est:yg44112.2 /CLONE=IMAGE:35277 /UG=Hs.224829 ESTs	R25692	1.06
KIAA1137 protein	AB032983	1.06
G protein-coupled receptor kinase 5	AI338653	1.06
melanoma antigen, family B, 2	NM_002364	1.06
NEL-like 1 (chicken)	NM_006157	1.06
kalikrein 2, prostatic	AA595465	1.06
testis enhanced gene transcript (BAX inhibitor 1)	NM_003217	1.06
related RAS viral (r-ras) oncogene homolog 2	AI753792	1.06
glypican 1	AI354864	1.06
hypothetical protein FLJ13491	NM_024623	1.06
hypothetical protein FLJ21616	NM_024587	1.06
aldehyde dehydrogenase 8 family, member A1	NM_002317	1.06
interactin 1 (SH3 domain protein)	NM_022568	1.06
core-binding factor, runt domain, alpha subunit 2; translocated to, 2	AW575493	1.06
porcupine	AF114488	1.06
secretory leukocyte protease inhibitor (antileukoprotease)	NM_005093	1.06
fertilization 1 homolog a	NM_022825	1.06
Consensus includes gb:BE857715 /FEA=EST /DB_XREF=gi:10372018 /DB_XREF=est:7g46g03.x1 /CLONE=IMAGE:3309556 /UG=Hs.8982 ESTs, Highly similar to KIAA1	NM_003084	1.06
DKFZP564M14.16 protein	AB62658	1.06
nuclear factor related to kappa B binding protein	BF981643	1.06
SRY (sex determining region Y)-box 30	AI887378	1.06
PR domain containing 2, with ZNF domain	NM_007017	1.06
nucleoporin 214KD (CAIN)	NM_012231	1.06
ecotrodes (multiple) 2	D14889	1.06
sperm associated antigen 6	AA188245	1.06
	AF078363	1.06

antigen identified by monoclonal antibodies 12E7, F21 and O13
Consensus includes gb:A1733515 /FEA=EST /DB_XREF=gi:5054676 /DB_XREF=est:ou99e05.x5 /CLONE=IMAGE:1635988 /UG=Hs.148907 Homo sapiens mRNA: cDNA
plectrophin (heparin binding growth factor 8, neurite growth-promoting factor 1)
kynureninase (L-tryptophan hydrolase)
peroxiredoxin 3
ubiquitin specific protease 14 (iRNA-guanine transglycosylase)
H4 histone family, member M
CED-6 protein
phosphoinositide-3-kinase, class 2, beta polypeptide
cytochrome c oxidase subunit VIa polypeptide 2
hypothetical protein DKFZp564L2423
alpha-methylacyl-CoA racemase
cyclin-dependent kinase 2
NADPH oxidase 1
TRF2-interacting telomeric RAP1 protein
RAS guanyl releasing protein 2 (calcium and DAG-regulated)
Wolf-Hirschhorn syndrome candidate 2
KIAA0144 gene product
book2 protein
chimerin (chimerin) 1
Consensus includes gb:AJ239383.1 /DEF=Homo sapiens mRNA for immunoglobulin heavy chain variable region, ID 31. /FEA=mRNA /GEN=IGHV /PROD=immunoglobulin
Consensus includes gb:A1357616 /FEA=EST /DB_XREF=gi:4109237 /DB_XREF=est:qu20g08.x1 /CLONE=IMAGE:1985374 /UG=Hs.101651 Homo sapiens mRNA: cDNA
collagen, type V, alpha 3
neural precursor cell expressed, developmentally down-regulated 4-like
relinitis pigmentosa GTPase regulator
nucleosomal binding protein 1
myosin IXB
glutamate receptor, metabotropic 2
elafin
serine/arginine repetitive matrix 1
RAD51 homolog (RecA homolog, E. coli) (S. cerevisiae)
cytochrome P450, subfamily I1C (mephenytoin 4-hydroxylase), polypeptide 18
KIAA0471 gene product
Consensus includes gb:A1561253 /FEA=EST /DB_XREF=gi:4511584 /DB_XREF=est:q27g06.x1 /CLONE=IMAGE:2210074 /UG=Hs.306649 Homo sapiens cDNA FLJ1340
hypothetical protein FLJ12998
5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-coupled)
KIAA0892 protein
serologically defined breast cancer antigen NY-BR-20
hypothetical protein MGC10798
cAMP responsive element binding protein 1
Consensus includes gb:AL036840 /FEA=EST /DB_XREF=gi:5408335 /DB_XREF=est:DKFZp564N1963_s1 /CLONE=DKFZp564N1963 /UG=Hs.22370 Homo sapiens mRNA
syndecan 2 (heparan sulfate proteoglycan 1, cell surface-associated, fibroglycan)
ribosomal protein L18
ras responsive element binding protein 1
hypothetical protein FLJ20654
ubiquitin specific protease 18
parvin, beta
lung type- cell membrane-associated glycoprotein
hypothetical protein FLJ20584
etoposide-induced mRNA
ubiquitously transcribed tetranucleotide repeat gene, X chromosome
hemochromatosis
KIAA1037 protein
GAS2-related on chromosome 22
Consensus includes gb:A875944 /FEA=EST /DB_XREF=gi:4650775 /DB_XREF=est:y65d07.x1 /CLONE=IMAGE:2283949 /UG=Hs.295963 ESTs
lipocalin 1 (protein migrating faster than albumin, tear prealbumin)
Toll-like receptor 8
cyclin-dependent kinase (CDC2-like) 10
guanine nucleotide binding protein (G protein), alpha activating activity polypeptide O
target of myo1-like 2 (chicken)
B1 for nudin
olfactory receptor, family 10, subfamily C, member 1
hypothetical protein MGC4276 similar to CG8198

hypothetical protein MGC2776
 chromodomain helicase DNA binding protein 1-like
 Consensus includes gb:BF347113 /FEA=EST /DB_XREF=gi:11294708 /DB_XREF=est:602020879F /CLONE=IMAGE:4156602 /UG=Hs.273138 ESTs
 KIAA1084 protein
 synaptogenin 2
 stomal antigen 3
 amiloride-sensitive cation channel 2, neuronal
 secretory granule, neuroendocrine protein 1 (7B2 protein)
 prostate epithelium-specific Ets transcription factor
 endothelial differentiation, sphingolipid G-protein-coupled receptor, 8
 collagen, type V, alpha 2
 KIAA0335 protein
 vsmn-like 1
 KIAA0185 protein
 apolipoprotein H (beta-2-glycoprotein I)
 ariadne homolog, ubiquitin-conjugating enzyme E2 binding protein, 1 (Drosophila)
 Consensus includes gb:AA293502 /FEA=EST /DB_XREF=gi:1941036 /DB_XREF=est:zt53n06.r1 /CLONE=IMAGE:726107 /UG=Hs.42733 CHMP1.5 protein /FL=gb:AF281
 ATP-binding cassette, sub-family C (CFTR/MRP), member 3
 caspase 2, apoptosis-related cysteine protease (neural precursor cell expressed, developmentally down-regulated 2)
 four and a half LIM domains 1
 hypothetical protein FLJ11004
 transforming growth factor beta-activated kinase-binding protein 1
 WNT1 inducible signaling pathway protein 1
 EGF-containing fibulin-like extracellular matrix protein 1
 dynein, axonemal, heavy polypeptide 9
 serine (or cysteine) proteinase inhibitor, clade F (alpha-2 antiplasmin, pigment epithelium derived factor), member 1
 HLA-B associated transcript 8
 Consensus includes gb:AW851066 /FEA=EST /DB_XREF=gi:7946583 /DB_XREF=est:IL3-CT0220-150200-070-802 /UG=Hs.314334 ESTs
 HIV-1 inducer of short transcripts binding protein: lymphoma related factor
 retinoid acid receptor, gamma
 potassium voltage-gated channel, shaker-related subfamily, beta member 1
 Consensus includes gb:AI097640 /FEA=EST /DB_XREF=gi:3445898 /DB_XREF=est:qb59a08.x1 /CLONE=IMAGE:1704374 /UG=Hs.66187 Homo sapiens done 23700 nt
 chromosome 6 open reading frame 32
 high-mobility group (nephilione chromosomal) protein 1
 tripartite motif-containing 14
 hypothetical protein FLJ13868
 KIAA0322 protein
 peptidylprolyl isomerase (cyclophilin)-like 2
 solute carrier family 21 (organic anion transporter), member 12
 nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor-like 1
 low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor)
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 8
 peptidylprolyl isomerase (cyclophilin)-like 2
 tissue factor pathway inhibitor (lipoprotein-associated coagulation inhibitor)
 ecdonucleotide pyrophosphatase/phosphodiesterase 1
 KIAA0582 protein
 Consensus includes gb:AI160126 /FEA=EST /DB_XREF=gi:3693506 /DB_XREF=est:qb51n08.x1 /CLONE=IMAGE:1703879 /UG=Hs.295901 KIAA0493 protein
 paired box gene 8
 tenascin XB
 hypothetical protein FLJ11328
 TAF4A RNA polymerase II, TATA box binding protein (TBP)-associated factor, 135 kD
 3-hydroxy-3-methylglutaryl-Coenzyme A synthase 2 (mitochondrial)
 hypothetical protein FLJ20559
 potassium inwardly-rectifying channel, subfamily J, inhibitor 1
 proline-rich protein BstII subfamily 1
 DnaJ (Hsp40) homolog, subfamily A, member 4
 Consensus includes gb:AL049991.1 /DEF=Homo sapiens mRNA; cDNA DKFZp564G222 (from done DKFZp564G222) /FEA=mRNA /DB_XREF=gi:4864243 /UG=Hs.2261
 meprin A, alpha (PABA peptide hydrolase)
 hypothetical protein DKFZp434P144
 killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 5
 fattyacyltransferase, CAAAX box, beta
 zinc finger RNA binding protein
 interferon, alpha 7

Consensus includes gb:AK021633.1 /DEF=Homo sapiens cDNA FLJ11571 fls, clone HEMBA1003328 /FEA=mRNA /DB_XREF=gi:10432853 /UG=Hs.306608 Homo sapiens AK021633
 Consensus includes gb:AK026449.1 /DEF=Homo sapiens cDNA FLJ22798 fls, clone KAA2544 /FEA=mRNA /DB_XREF=gi:10439319 /UG=Hs.308850 Homo sapiens cdi AK026449
 gb:NM_024118.1 /DEF=Homo sapiens hypothetical protein MGC4692 (MGC4692), mRNA /FEA=mRNA /GEN=MGC4692 /PROD=hypothetical protein MGC4692 /DB_XREF=gi:10439319 /UG=Hs.308850 Homo sapiens NM_024118
 Consensus includes gb:L23852.1 /DEF=Homo sapiens (clone Z146) retinal mRNA, 3' end and repeat region /FEA=mRNA /DB_XREF=gi:393126 /UG=Hs.73838 Homo sapiens L23852
 cyclin-dependent kinase inhibitor 1A (p21, Cip1)
 hypothetical protein FLJ13154
 Fanconi anemia, complementation group E
 Cluster incl. A057637.0ys1h06.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-1687483 /clone_end=3 /gb=A057637 /gi=3331503 /ug=Hs.21305 /len=598
 synaptosomal-associated protein, 29kD
 gb:NM_016421.1 /DEF=Homo sapiens relaxininsulin-like factor 1 (LOC51746), mRNA /FEA=cDS /GEN=LOC51746 /PROD=relaxininsulin-like factor 1 /DB_XREF=gi:7708 NM_016421
 KIAA0923 protein
 transcription factor similar to D. melanogaster homeodomain protein lady bird late
 hypothetical protein FLJ11222
 hypothetical protein FLJ14100
 eIF4E transporter
 T cell receptor gamma locus
 activating transcription factor 4 (tax-responsive enhancer element 867)
 heme oxygenase (decycling) 2
 SELENOPHOSPHATE SYNTHETASE : Human selenium donor protein
 serine/threonine kinase 38
 ubiquitin specific protease 4 (proto-oncogene)
 gb:AF095784.1 /DEF=Homo sapiens GABA-B receptor R2 (GABBR2) mRNA, complete cds /FEA=mRNA /GEN=GABBR2 /PROD=GABA-B receptor R2 /DB_XREF=gi:483 AF095784
 chorionic somatomammotropin hormone-like 1
 regulator of G-protein signalling 7
 neurofibromin 1 (neurofibromatosis, von Recklinghausen disease, Watson disease)
 hypothetical protein FLJ20894
 v-jun sarcoma virus 17 oncogene homolog (avian)
 Consensus includes gb:F1012 /FEA=EST /DB_XREF=gi:682615 /DB_XREF=est:HSC3AG012 /CLONE=c-3ag01 /UG=Hs.13252 Human EST clone 22453 mariner transposon F1012
 receptor-associated protein of the synapse, 43kD
 KIAA0878 protein
 solute carrier family 17 (sodium phosphate), member 4
 hypothetical protein FLJ21859
 HSPC133 protein
 solute carrier family 16 (monocarboxylic acid transporters), member 5
 CBF1 interacting corepressor
 sarcosine dehydrogenase
 bridging integrator 1
 Consensus includes gb:AW238724 /FEA=EST /DB_XREF=gi:6571113 /DB_XREF=est:sp30h08.x1 /CLONE=IMAGE-2741919 /UG=Hs.314230 ESTs, Highly similar to clock AW238724
 centromere protein F (350/400kD, mitotin)
 poliovirus receptor
 hypothetical protein
 colony stimulating factor 1 (macrophage)
 putative UDP-GalNAc:polypeptide N-acetylglucosaminyltransferase T9
 lymphocyte antigen 84 homolog, radioprotective 105kD (mouse)
 oral-facial-digital syndrome 1
 Hermansky-Pudlak syndrome
 protein phosphatase 1, regulatory subunit 10
 transmembrane protease, serine 3
 dystrophin (muscular dystrophy, Duchenne and Becker types), includes DXS142, DXS164, DXS208, DXS230, DXS238, DXS269, DXS270, DXS272
 hypothetical protein FLJ22593
 integrin, alpha 7
 G6C protein
 interferon (alpha, beta and omega) receptor 1
 cell growth regulatory with ring finger domain
 Fc fragment of IgG, low affinity IIa, receptor for (CD32)
 hypothetical protein, estradiol-induced
 interleukin 15
 single-stranded DNA-binding protein 1
 KIAA0527 protein
 inhibin, beta C
 ATP-binding cassette, sub-family C (CFTR/MRP), member 3
 myobulbin related protein 8
 hypothetical protein FLJ23151

hypothetical protein FLJ21802
 a disintegrin and metalloproteinase domain 20
 leptin (murine obesity homolog)
 LIM domain binding 3
 KIAA0728 protein
 DnaJ (Hsp40) homolog, subfamily B, member 12
 cofactor required for Sp1 transcriptional activation, subunit 2 (150KD)
 granzyme A (granzyme 1, cytotoxic T-lymphocyte-associated serine esterase 3)
 hypothetical protein P1 p373c6
 hypothetical protein FLJ23323
 rhodopsin (opsin 2, rod pigment) (retinitis pigmentosa 4, autosomal dominant)
 Consensus includes gb:AK026465.1 /DEF=Homo sapiens cDNA: FLJ22812 fls, clone KIAU2955, /FEA=mRNA /DB_XREF=gj:0439335 /UG=Hs.306853 Homo sapiens cdi AK026465
 hypothetical protein MGC14797
 homolog of mouse C2PA
 tripartite motif-containing 2
 Consensus includes gb:AW451806 /FEA=EST /DB_XREF=gj:5992582 /DB_XREF=est:U1-H-B13-alk-c-10-Q-U1.s1 /CLONE=IMAGE:2737146 /UG=Hs.32458 ESTs
 5-hydroxytryptamine (serotonin) receptor 3A
 secreted protein of unknown function
 solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 6
 lymphocyte antigen 117
 olfactory receptor, family 7, subfamily E, member 47 pseudogene
 gb:BC000233.1 /DEF=Homo sapiens, Similar to guanine nucleotide binding protein, alpha transducing 2, clone MGC:2232, mRNA, complete cds, /FEA=mRNA /PROD=Sin
 ADP-ribosylation factor-like 7
 Consensus includes gb:AK026778.1 /DEF=Homo sapiens cDNA: FLJ23125 fls, clone LNG08217, /FEA=mRNA /DB_XREF=gj:10439709 /UG=Hs.78325 step II splicing fact
 serum amyloid A2
 follistatin
 heparin-binding (tenascin C, cytotactin)
 Consensus includes gb:AV661514 /FEA=EST /DB_XREF=gj:3882528 /DB_XREF=est:AV661514 /CLONE=GLCGTC04 /UG=Hs.282693 ESTs
 synaptosomal-associated protein, 23kD
 much 4, tracheobronchial
 KIAA0077 protein
 seroglycan, beta (43kD dystrophin-associated glycoprotein)
 hypothetical protein PRO1866
 UDP-Gal4betaGalNAc beta 1,4'-galactosyltransferase, polypeptide 1
 ADAM-like, decalin 1
 interferon, gamma-inducible protein 16
 Consensus includes gb:BF542055 /FEA=EST /DB_XREF=gj:11629436 /DB_XREF=est:602068320F1 /CLONE=IMAGE:4067357 /UG=Hs.190785 ESTs
 calcium channel, voltage-dependent L type, alpha 1S subunit
 5'-nucleotidase (purine), cytosolic type B
 vascular Rab-GAP78C-containing
 hypothetical protein FLJ20772
 likely ortholog of mouse D11gpr2
 protective protein for beta-galactosidase (galactosialcosis)
 Consensus includes gb:AK021533.1 /DEF=Homo sapiens cDNA FLJ11471 fls, clone HEMBA1001675, weakly similar to VACUOLAR PROTEIN SORTING-ASSOCIATED P
 CASP8 and FADD-like apoptosis regulator
 nucleoporin 214KD (CAIN)
 ribosomal protein L14
 TONDU
 tryptase beta 2
 glyceraldehyde-3-phosphate dehydrogenase, testis-specific
 aconitase 2, mitochondrial
 melanin, ring finger protein, 3
 leucine zipper, putative tumor suppressor 1
 leukocyte-associated Ig-like receptor 1
 guanine nucleotide binding protein (G protein), beta polypeptide 3
 Consensus includes gb:AK025041.1 /DEF=Homo sapiens cDNA: FLJ21388 fls, clone COL03424, /FEA=mRNA /DB_XREF=gj:0437478 /UG=Hs.306751 Homo sapiens cC AK025041
 Consensus includes gb:AT39005 /FEA=EST /DB_XREF=gj:5100986 /DB_XREF=est:w34b03.x1 /CLONE=IMAGE:2392109 /UG=Hs.173448 KIAA0418 gene product
 hypothetical protein FLJ00002
 protein arginine N-methyltransferase 3(hnRNP methyltransferase S. cerevisiae)-like 3
 testis-specific kinase substrate
 Consensus includes gb:AK026889.1 /DEF=Homo sapiens cDNA: FLJ23236 fls, clone COL00725, /FEA=mRNA /DB_XREF=gj:0439654 /UG=Hs.288786 Homo sapiens cC AK026889
 pre-mRNA splicing factor 17
 BG528818

Consensus includes gb:AL049988.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564F212 (from clone DKFZp564F212). /FEA=mRNA /DB_XREF=gi:4884239 /UG=Hs.30631 AL049988
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 4
eukaryotic translation elongation factor 1 epsilon 1
eyes absent homolog 4 (Drosophila)
Consensus includes gb:AK025114.1 /DEF=Homo sapiens cDNA: FLJ21481 fls, clone COL04727. /FEA=mRNA /DB_XREF=gi:10437566 /UG=Hs.306768 Homo sapiens cDNA
DKFZP434N014 protein
silver homolog (mouse)
KIAA0633 protein
cell division cycle 27
DKFZP586N0618 protein
KIAA0893 protein
methylenetetrahydrofolate phosphorylase
leptin receptor
Consensus includes gb:AL137435.1 /DEF=Homo sapiens mRNA: cDNA DKFZp761G0924 (from clone DKFZp761G0924). /FEA=mRNA /DB_XREF=gi:6807994 /UG=Hs.27 AL137435
complement component 5
KIAA0665 gene product
Consensus includes gb:AK024548.1 /DEF=Homo sapiens cDNA: FLJ20895 fls, clone ADK4Q3483. /FEA=mRNA /DB_XREF=gi:10436853 /UG=Hs.285737 Homo sapiens cDNA
transmembrane gamma-carboxyglutamic acid protein 3
latent transforming growth factor beta binding protein 4
Chediak-Higashi syndrome 1
linker for activation of T cells
hypothetical protein PRO1316
centromere protein E (312kD)
fibroblast growth factor 21
sulfotransferase family, cytosolic, 1B, member 1
ribonuclease P (30kD)
engrailed homolog 2
KIAA1053 protein
H4 histone family, member H
likely homolog of mouse immunity-associated nucleotide 4
potassium large conductance calcium-activated channel, subfamily M, alpha member 1
hypothetical protein MGCS627
hydroxyprostaglandin dehydrogenase 15-(NAD)
Consensus includes gb:N35922 /FEA=EST /DB_XREF=gi:1157084 /DB_XREF=est:Y28905.s1 /CLONE=IMAGE:272600 /UG=Hs.269852 ESTs. Weakly similar to ALU1_H N35922
Consensus includes gb:AL046979 /FEA=EST /DB_XREF=gi:5435035 /DB_XREF=est:DKFZp586K0617 /UG=Hs.9973 tennin
neuregulin 1
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 3
protease, serine, 1 (trypsin 1)
Consensus includes gb:AU147698 /FEA=EST /DB_XREF=gi:11009219 /DB_XREF=est:AU147698 /CLONE=MAMMA1001408 /UG=Hs.306833 Homo sapiens cDNA FLJ12
Consensus includes gb:BF131965 /FEA=EST /DB_XREF=gi:10971005 /DB_XREF=est:601820948F1 /CLONE=IMAGE:4052867 /UG=Hs.10684 Homo sapiens clone 2442 BF131965
TIA1 cytotoxic granule-associated RNA binding protein
hypothetical protein FLJ13158
homolog of mouse BMP-2 inducible kinase
hypothetical protein
Consensus includes gb:AA114843 /FEA=EST /DB_XREF=gi:1669911 /DB_XREF=est:20212.s1 /CLONE=IMAGE:491183 /UG=Hs.183454 Homo sapiens mRNA: cDNA DI AA114843
myelin basic protein
lymphoid enhancer-binding factor 1
cellular zinc finger anti-NF-kappaB Cezanne
desmoglein 1
a disintegrin-like and metalloprotease (repolysin type) with thrombospondin type 1 motif, 2
chorionic gonadotropin hormone 2
taste receptor, type 2, member 3
sialidase 2 (cytosolic sialidase)
SKIP (for skeletal muscle and kidney enriched inositol phosphatase)
splicing factor, arginine/serine-rich 11
B-cell CLL lymphoma 11A (zinc finger protein)
Consensus includes gb:AL049279.1 /DEF=Homo sapiens mRNA: cDNA DKFZp564I083 (from clone DKFZp564I083). /FEA=mRNA /DB_XREF=gi:4500035 /UG=Hs.202886 AL049279
Cdc42 effector protein 3
Rag D protein
protocadherin alpha 8
found in inflammatory zone 3
chromosome 14 open reading frame 2
transcriptional adaptor 2 (ADA2 homolog, yeast)-like

Integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)
 runt-related transcription factor 1 (acute myeloid leukemia 1; arn1 oncogene)
 melanoma-associated chondroitin sulfate proteoglycan-like
 protein tyrosine phosphatase, non-receptor type substrate 1
 tumor susceptibility gene 101
 protein tyrosine phosphatase, receptor-type, Z polypeptide 1
 enhancer of zeste homolog 1 (Drosophila)
 KIAA0391 gene product
 tumor necrosis factor receptor superfamily, member 5
 ketch-like 3 (Drosophila)
 gb:M33376.1/DEF=Human pseudo-chloride reductase mRNA, complete cds. /FEA=mRNA /DB_XREF=gi:187444 /FL=gb:M33376.1
 WD40 repeat domain 11 protein
 KIAA1105 protein
 transcription factor AP-2, alpha (activating enhancer binding protein 2 alpha)
 cathepsin O
 hyperpolarization activated cyclic nucleotide-gated potassium channel 2
 fibroblast growth factor 18
 Consensus includes gb:AL050065.1 /DEF=Homo sapiens mRNA; cDNA DKFZp566M043 (from clone DKFZp566M043). /FEA=mRNA /DB_XREF=gi:4884295 /UG=Hs.2121 AL050065
 hypothetical protein FLJ20151
 CK2 interacting protein 1; HQ0024c protein
 HBST-like (S. cerevisiae)
 hypothetical protein FLJ13102
 S100 calcium binding protein A10 (annexin II ligand, calpactin I, light polypeptide (p11))
 hypothetical protein MG3113
 SH2-B homolog
 activated p21cdc42Hs kinase
 regulator of mitotic spindle assembly 1
 calcium/calmodulin-dependent protein kinase (CaM kinase) II beta
 hypothetical protein FLJ10696
 aquaporin 3
 muscle RAS oncogene homolog
 Consensus includes gb:AI206718 /FEA=EST /DB_XREF=gi:3765390 /DB_XREF=est:qf61b12.x1 /CLONE=IMAGE:1754495 /UG=Hs.143920 Homo sapiens DNA from chro
 MAD, mothers against decapentaplegic homolog 3 (Drosophila)
 myotubularin related protein 6
 sirin, silent mating type information regulation 2 homolog 3 (S. cerevisiae)
 MAX dimerization protein
 hypothetical protein FLJ10232
 chromosome 9 open reading frame 3
 Huntingtin interacting protein H
 collagen, type XIX, alpha 1
 KIAA1243 protein
 dual specificity phosphatase 8
 hepatocellular carcinoma-associated antigen 112
 Consensus includes gb:BF593594 /FEA=EST /DB_XREF=gi:11885918 /DB_XREF=est:7q12a08.x1 /CLONE=IMAGE:3697838 /UG=Hs.274440 Homo sapiens mRNA; cDN
 chaperonin containing TCP1, subunit 8A (zeta 1)
 Consensus includes gb:AL031651 /DEF=Human DNA sequence from clone RP5-1054A22 on chromosome 20q11.22-12 Contains two isoforms of the gene for TGM2 (trans
 nuclear receptor subfamily 2, group C, member 1
 integrin, alpha 6
 calicin
 Cluster Inc. AB970898:wr1cd3.x1 Homo sapiens cDNA, 3' end /clone=IMAGE:2488324 /clone_end=3 /gb=AB970898 /gi=57877724 /ug=Hs.234898 /len=382
 thoredoxin-like, 32KD
 olfactory receptor, family 7, subfamily E, member 12 pseudogene
 single-minded homolog 2 (Drosophila)
 adenyate cyclase 6
 cyclic AMP phosphoprotein, 19 kD
 Consensus includes gb:AB79213 /FEA=EST /DB_XREF=gi:4869395 /DB_XREF=est:u72p04.x1 /CLONE=IMAGE:2256630 /UG=Hs.81920 ESTs. Weakly similar to cDNA I
 BTB and CNC homology 1, basic leucine zipper transcription factor 1
 hypothetical protein
 hypothetical protein FLJ10055
 carnitine palmitoyltransferase 1, liver
 transcription factor Dp-2 (E2F dimerization partner 2)
 paired basic amino acid cleaving system 4
 hypothetical protein FLJ10738

baculoviral IAP repeat-containing 4
 immunoglobulin lambda variable (V)OR22-1
 hypothetical protein FLJ22233
 pre-mRNA processing factor 31 homolog (yeast)
 gb:M32577.1 /DEF=Human MHC HLA-DQ beta mRNA, complete cds. /FEA=mRNA /GEN=HLA-DQB1 /DB_XREF=gi:188194 /FL=gb:M32577.1
 DKFZP564B147 protein
 NIF3 NGG1 interacting factor 3-like 1 (S. pombe)
 C-type (calcium dependent, carbohydrate-recognition domain) lectin, superfamily member 2 (activation-induced)
 hypothetical protein DKFZp434L0718
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 5
 cAMP responsive element modulator
 stannocalcin 1
 diacylglycerol kinase, alpha (80kD)
 hypothetical protein FLJ10349
 keratin 6A
 Interleukin 1 receptor accessory protein
 (clone PWHL22-24) myosin light chain 2
 hypothetical protein FLJ10388
 endometrial bleeding associated factor (left-right determination), factor A; transforming growth factor beta superfamily)
 transforming, acidic coiled-coil containing protein 3
 tropomyosin 2 (beta)
 Consensus includes gb:AL157388 /DEF=Human DNA sequence from clone RP11-56H7 on chromosome 10. Contains ESTs, STSs and GSSs. Contains the gene for the ne
 HIV-1 Rev binding protein
 hypothetical protein FLJ10856
 von Willebrand factor
 Consensus includes gb:AU145354 /FEA=EST /DB_XREF=gi:11006875 /DB_XREF=est:AU145354 /CLONE=HEMBA1004606 /UG=Hs.287411 Homo sapiens cDNA FLJ111
 KIAA0097 protein
 mannan-binding lectin serine protease 1 (CA/C2 activating component of Rf-reactive factor)
 CCAAT/enhancer binding protein (C/EBP), alpha
 zinc finger protein 148 (pH2-52)
 ELK1, member of ETS oncogene family
 KIAA0248 protein
 Ras-related GTP-binding protein
 son of sevenless (Drosophila) homolog 2
 Cluster incl. AA524093.ng3304.s1 Homo sapiens cDNA, 3' end /clone=IMAGE:936607 /clone_end=3' /gb=AA524093 /gi=2265021 /ug=Hs.23158 /len=718
 erythropoietin receptor
 H1 histone family, member 0
 glycerol kinase
 solute carrier family 12 (sodium/potassium/chloride transporters), member 2
 activating transcription factor 2
 cystinosis, nephropathic
 transcription factor-like 1
 ribonuclease, RNase A family, 4
 hypothetical protein MGC2849
 tyrosinase beta 1
 natriuretic peptide precursor B
 hypothetical protein
 RNA binding motif protein 9
 ems1 sequence (mammary tumor and squamous cell carcinoma-associated (p80/85 src substrate)
 insulin-like growth factor 2 receptor
 Consensus includes gb:AK024525.1 /DEF=Homo sapiens cDNA: FLJ20872 fts, clone AOKA02604. /FEA=mRNA /DB_XREF=gi:10438827 /UG=Hs.306683 Homo sapiens
 neuromedin B
 protein phosphatase 2A, regulatory subunit B' (PR 53)
 fts485
 karyopherin beta 2b, transportin
 hypothetical protein FLJ10579
 ubiquitin activating enzyme E1-like protein
 nescent helix loop helix 2
 gap junction protein, alpha 3, -46kD (connexin 46)
 hypothetical protein FLJ13188
 eukaryotic translation initiation factor 4E-like 3
 Consensus includes gb:AL050053.1 /DEF=Homo sapiens mRNA: cDNA DKFZp566E103 (from clone DKFZp566E103) /FEA=mRNA /DB_XREF=gi:4864289 /UG=Hs.2273:AL050053
 KIAA0793 gene product

ataxin 2 related protein	NM_007245	1.04
KIAA1882 protein	BC003650	1.04
SIPL protein	NM_018269	1.04
cydn D2	AW026491	1.04
hypothetical protein FLJ22187	NM_024533	1.04
hypothetical protein DKFZp434A1022	NM_021165	1.04
histone acetyltransferase 1	NM_003642	1.04
epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogene homolog, avian)	NM_005228	1.04
Consensus includes gb:R61519 /FEA=EST /DB_XREF=gi:832214 /DB_XREF=est:YH1804.s1 /CLONE=IMAGE:37896 /UG=Hs.185807 Homo sapiens clone 24758 mRNA	R61519	1.04
nuclear factor of activated T-cells, cytoplasmic, calcineurin-dependent 1	U08015	1.04
calcium/calmodulin-dependent protein kinase kinase 2, beta	AA181179	1.04
erythropoietin receptor	M60459	1.04
modlin	NM_002418	1.04
1003 protein	NM_002322	1.04
disabled homolog 2, mitogen-responsive phosphoprotein (Drosophila)	AF188296	1.04
myosin, light polypeptide 3, alkali; ventricular, skeletal, slow	NM_000256	1.04
cytochrome P450 polypeptide 43	AF280110	1.04
ets variant gene 5 (ets-related molecule)	NM_004454	1.04
cadherin 7, type 2	NM_004381	1.04
glutathione S-transferase A3	NM_000847	1.04
Consensus includes gb:AL080182.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434O151 (from clone DKFZp434O151) /FEA=mRNA /DB_XREF=gi:52626568 /UG=Hs.2251	AL080182	1.04
a disintegrin and metalloproteinase domain 17 (tumor necrosis factor, alpha, converting enzyme)	U86755	1.04
plasminogen activator, urokinase receptor	AY029180	1.04
chloride channel 3	AI760629	1.04
hypothetical protein FLJ20707	NM_017936	1.04
UV-B repressed sequence, HUR 7	BE146534	1.04
glutathione S-transferase subunit 13 homolog	NM_015917	1.04
clone FLB3816	NM_016415	1.04
KIAA1048 protein	AW451954	1.04
serologically defined colon cancer antigen 16	NM_008649	1.04
zinc finger protein, subfamily 1A, 2 (Helios)	NM_018260	1.04
monamine oxidase A	AA923354	1.04
hypothetical protein MGC5356	NM_024059	1.04
PROO214 protein	NM_014120	1.04
Consensus includes gb:AK024873.1 /DEF=Homo sapiens cDNA; FLJ21220 fts, clone COL00546 /FEA=mRNA /DB_XREF=gi:10437285 /UG=Hs.306713 Homo sapiens cC	AK024873	1.04
Fancconi anemia, complementation group C	NM_000136	1.04
transcription factor 8 (represses interleukin 2 expression)	AI373166	1.04
hypothetical protein FLJ22175	AA551370	1.04
nuclear mitotic apparatus protein 1	AI337584	1.04
hypothetical protein MGC3123	NM_024107	1.04
KIAA0982 protein	AK022530	1.04
paralemmin	NM_002579	1.04
hypothetical protein FLJ12838	NM_024641	1.04
dactylund homolog (Drosophila)	NM_004392	1.04
hypothetical protein FLJ23229	NM_025058	1.04
insulin-like growth factor 1 (somatomedin C)	MZ9644	1.04
adenosine A2b receptor pseudogene	AL022398	1.04
N-myc downstream-regulated gene 3	BC001805	1.04
Consensus includes gb:BG150564 /FEA=EST /DB_XREF=gi:12662594 /DB_XREF=est:702g04.x1 /CLONE=IMAGE:3443143 /UG=Hs.184326 CDC10 (cell division cycle	BG150564	1.04
inosine triphosphatase (nucleoside triphosphate pyrophosphatase)	AF219116	1.04
phenylalanine hydroxylase	NM_000277	1.04
calcium channel, voltage-dependent, alpha 2/delta 3 subunit	NM_016398	1.04
KIAA0576 protein	AU144775	1.04
hypothetical protein	NM_019073	1.04
adaptor-related protein complex 1, sigma 1 subunit	NM_001283	1.04
small inducible cytokine subfamily A (Cys-Cys), member 11 (eotaxin)	D46372	1.04
formyltetrahydrofolate dehydrogenase	NM_012190	1.04
EphB2	D31681	1.04
progesterone receptor	NM_000926	1.04
paired-like homeodomain transcription factor 3	NM_005029	1.04
phosphothiosulfate-3-kinase, regulatory subunit, polypeptide 3 (p55, gamma)	AF028765	1.04
hypothetical protein FLJ11618	NM_022452	1.04
zinc finger protein 238	NM_006352	1.04

KIAA0427 gene product	U55962	1.04
leukocyte-associated Ig-like receptor 1	NM_021708	1.04
KIAA0261 protein	AK027005	1.04
hypothetical protein	NM_015689	1.04
TGF-beta-induced factor 2 (TALE family homeobox)	NM_021809	1.04
bladder cancer associated protein	NM_006898	1.04
hypothetical protein FLJ11585	AI743854	1.04
glutamate receptor, ionotropic, N-methyl D-aspartate 2D	NM_000836	1.04
DKFZ586P2219 protein	W80642	1.04
KIAA0717 protein	AY008093	1.04
endocrine regulator	AW613549	1.04
nuclear autoantigen (55kD) similar to rat synaptonemal complex protein	NM_006455	1.04
aldohyde dehydrogenase 1 family, member A1	NM_000689	1.04
elongation of very chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1	NM_016031	1.04
homolog of yeast long chain polyunsaturated fatty acid elongation enzyme 2	AL136939	1.04
kinesin family member 4A	NM_012310	1.04
tol-like receptor 4	NM_003266	1.04
hypothetical protein FLJ20184	NM_017700	1.04
exosome component Rps41	NM_018037	1.04
hypothetical protein FLJ20279	NM_017740	1.04
Consensus includes gb:AB007970.1 /DEF=Homo sapiens mRNA, chromosome 1 specific transcript KIAA0501. /FEA=mRNA /DB_XREF=gi:3413945 /UG=Hs.223020 Homi	AB007970	1.04
chorionic gonadotropin hormone 1 (placental lactogen)	BC005921	1.04
signal transducer and activator of transcription 5B	NM_012448	1.04
hypothetical protein FLJ21007	NM_030794	1.04
Consensus includes gb:AI695595 /FEA=EST /DB_XREF=gi:4983495 /DB_XREF=est:we49g06.x1 /CLONE=IMAGE:2344474 /UG=Hs.293219 ESTs	AI695595	1.04
coagulation factor III (thromboplastin, tissue factor)	NM_001993	1.04
proliferase bogenesis factor 13	NM_002918	1.04
Consensus includes gb:AK024177.1 /DEF=Homo sapiens cDNA FLJ14115 fis, clone MAMMA1001760. /FEA=mRNA /DB_XREF=gi:10436493 /UG=Hs.266263 Homo sapie	AK024177	1.04
hypothetical protein 24636	AI150186	1.04
promyelocytic leukemia	M78462	1.04
Consensus includes gb:L34409.1 /DEF=Homo Sapiens (clone B383E13) chromosome 4p16.3 DNA fragment. /FEA=mRNA /DB_XREF=gi:800517 /UG=Hs.94789 Homo S	L34409	1.04
proline arginine-rich and leucine-rich repeat protein	NM_002725	1.04
MAD2 mitotic arrest deficient-like 1 (yeast)	NM_002358	1.04
putative DNA binding protein	AF072814	1.04
lysine aminotransferase	AV647713	1.04
KIAA0342 gene product	AA035414	1.04
wingless-type MMTV integration site family, member 16	NM_016087	1.04
MAD, mothers against decapentaplegic homolog 3 (Drosophila)	NM_005802	1.04
growth suppressor 1	NM_022356	1.04
KIAA1087 protein	AB028980	1.04
cytochrome P450, subfamily IIIA (naphthalene oxidase), polypeptide 5	NM_000777	1.04
promyelocytic leukemia	NM_002675	1.04
G protein-coupled receptor kinase 6	AF040752	1.04
N-acetylglucosamine amidohydrolase (acid ceramidase)	AI934569	1.04
Consensus includes gb:AL133519 /DEF=Human DNA sequence from clone RP11-12201 on chromosome 20 Contains a novel gene encoding a protein orthologous to the	AL133519	1.04
PKS1 postmitotic segregation increased 1 (S. cerevisiae)	BG434893	1.04
Consensus includes gb:AW081113 /FEA=EST /DB_XREF=gi:6036265 /DB_XREF=est:xc29c08.x1 /CLONE=IMAGE:2585678 /UG=Hs.18368 DKFZP56480769 protein	AW081113	1.04
CGI-86 protein	AF126782	1.04
mitogen-activated protein kinase kinase kinase 4	NM_004834	1.04
small nuclear ribonucleoprotein polypeptide E	NM_003094	1.04
Consensus includes gb:AF005082.1 /DEF=Homo sapiens skin-specific protein (xp33) mRNA, partial cds. /FEA=mRNA /GEN=xp33 /PROD=skin-specific protein /DB_XREF=AF005082	AF005082	1.04
hypothetical protein p5326	AF073483	1.04
cyclin-dependent kinase 8	R59697	1.04
KIAA1074 protein	NM_014915	1.04
dual-specificity tyrosine-(Y)-phosphorylation regulated kinase 2	Y09216	1.04
E74-like factor 2 (eta domain transcription factor)	NM_006874	1.04
tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain	AF012538	1.04
block of proliferation 1	AK024840	1.04
Consensus includes gb:AL137325.1 /DEF=Homo sapiens mRNA; cDNA DKFZp434M0835 (from clone DKFZp434M0835) /FEA=mRNA /DB_XREF=gi:6607816 /UG=Hs.27 AL137325	AL137325	1.04
SON DNA binding protein	NM_003103	1.04
basic helix-loop-helix domain containing, class B, 2	BG326045	1.04
uncharacterized hypothalamus protein HT012	NM_018473	1.04
nidogen 2	NM_007361	1.04

kalikrein 2, prostatic
putative nucleotide binding protein, estradiol-induced
Consensus includes gb:M28219.1 /DEF=Homo sapiens low density lipoprotein receptor (FH 10 mutant causing familial hypercholesterolemia) mRNA, 3 end. /FEA=mRNA / M28219
Consensus includes gb:AU144294 /FEA=EST /DB_XREF=gi:1005815 /DB_XREF=est:AU144294 /CLONE=HEMBA1001463 /UG=Hs.288530 Homo sapiens cDNA FLJ11, AU144294
Consensus includes gb:AL133604.1 /DEF=Homo sapiens mRNA, cDNA DKFZp434G1615 (from clone DKFZp434G1615). /FEA=mRNA /DB_XREF=gi:6598212 /UG=Hs.27 AL133604
hypothetical protein FLJ22501
Tax1 (human T-cell leukemia virus type 1) binding protein 1
Consensus includes gb:AA912711 /FEA=EST /DB_XREF=gi:3052103 /DB_XREF=est:q30108.s1 /CLONE=IMAGE:1524999 /UG=Hs.26395 erythrocyte membrane protein 1 AA912711
hypothetical protein DKFZp667O2416
Consensus includes gb:AI634532 /FEA=EST /DB_XREF=gi:4685862 /DB_XREF=est:bt52a09.x1 /CLONE=IMAGE:2273224 /UG=Hs.13423 Homo sapiens clone 24468 mR AI634532
siyltransferase 4B (beta-galactosidase alpha-2,3-syltransferase)
acid phosphatase 2, lysosomal
melanophilin
KJAA1034 protein
melanoma cell adhesion molecule
reserved
splicing factor 3a, subunit 1, 120KD
loss of heterozygosity, 11, chromosomal region 2, gene A
cytochrome P450, subfamily 11A (phenobarbital-inducible), polypeptide 7, pseudogene 1
GM2 ganglioside activator protein
early growth response 4
myoglobin
calponin 2
WD repeat domain 10
I-box and leucine-rich repeat protein 7
5-hydroxytryptamine (serotonin) receptor 4
hypothetical protein FLJ10698
terazspan 5
cholinergic receptor, nicotinic, beta polypeptide 2 (neuronal)
hypothetical protein FLJ10469
putative selenocysteine lyase
transcription factor-like 4
ATPase, Na+/K+ transporting, beta 1 polypeptide
WW domain-containing protein 1
hypothetical protein FLJ10600
membrane-bound transcription factor protease, site 2
hypothetical protein FLJ20417
KIAA0626 gene product
zinc finger protein 154 (pH2-92)
utrophin (homologous to dystrophin)
hepatocyte growth factor (hepatopoietin A, scatter factor)
zinc finger protein 304
Consensus includes gb:AC003989 /DEF=Human BAC clone CTB-7.J15 from Tq31 /FEA=CDS /DB_XREF=gi:2772538 /UG=Hs.24069 Human BAC clone CTB-7.J15 from 7 AC003989
hypothetical protein FLJ12886
pregnancy specific beta-1-glycoprotein 7
Snf2-related CBP activator protein
B-cell CLL/lymphoma 3
transcription factor 21
tweety homolog 1 (Drosophila)
potassium voltage-gated channel, shaker-related subfamily, member 3
ubiquitin specific protease 2
DKFZP564M1415 protein
small nuclear ribonucleoprotein D1 polypeptide (16KD)
chymotrypsin-like
GATA binding protein 2
Consensus includes gb:AI274196 /FEA=EST /DB_XREF=gi:3896464 /DB_XREF=est:q46b12.x1 /CLONE=IMAGE:1875359 /UG=Hs.292368 ESTs
inositol 1,4,5-trisphosphate 3-kinase B
chorionic somatomammotropin hormone-like 1
sarcoma amplified sequence
DEAD/HD (Asp-Glu-Ala-Asp/His) box polypeptide 24
synaptophysin 1
retinoblastoma 1 (including osteosarcoma)
adrenergic, beta, receptor kinase 1

BC005196 1.04
NM_014366 1.04
M28219 1.04
AU144294 1.04
AL133604 1.04
NM_024747 1.04
AF090891 1.04
AA912711 1.04
BG168720 1.04
AI634532 1.04
NM_006927 1.04
NM_001610 1.04
NM_024101 1.04
AB028957 1.04
BC006329 1.04
NM_020646 1.04
NM_005877 1.04
BC001234 1.04
U22830 1.04
X61094 1.04
NM_001965 1.04
NM_005368 1.04
NM_004368 1.04
AK000773 1.04
AU145127 1.04
NM_000870 1.04
AL050219 1.04
AF065389 1.04
NM_000748 1.04
NM_018102 1.04
AA911739 1.04
N40555 1.04
NM_001677 1.04
BF131791 1.04
NM_018152 1.04
U73479 1.04
NM_017810 1.04
AF327560 1.04
U20648 1.04
NM_007124 1.04
M77227 1.04
NM_020857 1.04
AC003989 1.04
NM_019108 1.04
NM_002783 1.04
AB002307 1.04
AI829875 1.04
NM_003206 1.04
NM_020659 1.04
NM_002232 1.04
NM_004205 1.04
BF966021 1.04
NM_006938 1.04
BF506665 1.04
NM_002050 1.04
AI274196 1.04
NM_002221 1.04
NM_022580 1.04
AL514076 1.04
NM_020414 1.04
BF511590 1.04
M19701 1.04
U08438 1.04

cystinosis, nephropathic	NM_004937	1.04
zodlin related factor 1	AI338837	1.04
secretory pathway component Sec31B-1	AF274863	1.04
Neuro-d4 (rat) homolog	NM_004647	1.04
myosin, light polypeptide 1, alkali; skeletal, fast	M20643	1.04
fatty acid amide hydrolase	NM_001441	1.04
protein similar to E.coli yhdg and R. capsulatus nifR3	NM_007016	1.04
lipase protein	AF225418	1.04
glutamate receptor, ionotropic, AMPA 4	NM_000829	1.04
ligase III, DNA, ATP-dependent	NM_013975	1.04
hypothetical protein MGC2745	NM_024117	1.04
collagen, type VI, alpha 2	AL531750	1.04
regulator of G-protein signalling 1	SS9049	1.04
Consensus includes gb:AW663883 /FEA=EST /DB_XREF=gi:7456424 /DB_CLONE=IMAGE:2977699 /UG=Hs.43834 Homo sapiens immunoglobulin	AF663885	1.04
briefdin A-inhibited guanine nucleotide-exchange protein 1	AF084520	1.04
calcium channel, voltage-dependent, gamma subunit 2	AL022313	1.04
major histocompatibility complex, class II, DR beta 4	BC005312	1.04
fused leuc homolog (mouse)	NM_022476	1.04
nuclear receptor coactivator 3	AF010227	1.04
calcineurin binding protein 1	NM_012295	1.04
prostate epithelium-specific Ets transcription factor	NM_012391	1.04
acrosomal vesicle protein 1	NM_020113	1.04
potassium voltage-gated channel, subfamily H (eag-related), member 2	AB044806	1.04
dual specificity phosphatase 7	X93921	1.04
peroxisome proliferative activated receptor, delta	L07592	1.04
tumor necrosis factor receptor superfamily, member 6b, decoy	NM_003823	1.04
absent in melanoma 1	U83115	1.04
vesicle-associated membrane protein 5 (myobrevin)	AB14468	1.04
zinc finger protein 202	N81520	1.04
serine/threonine kinase 22B (spemiotogenesis associated)	AB52441	1.04
tousted-like kinase 1	NM_012290	1.04
Consensus includes gb:AW517688 /FEA=EST /DB_XREF=gi:7155768 /DB_CLONE=IMAGE:2776521 /UG=Hs.305923 ATPase, Ca++ transporting,	AW517686	1.04
nodulin 2	NM_004475	1.04
makorin, ring finger protein, 1	NM_013448	1.04
myobulbin related protein 2	NM_016156	1.04
potassium voltage-gated channel, Isk-related family, member 1	NM_000219	1.04
hypothetical protein FLJ20342	NM_017774	1.04
hypothetical protein FLJ22729	NM_024683	1.04
polymerase (DNA directed), gamma 2, accessory subunit	NM_007215	1.04
KIAA0980 protein	NM_025176	1.04
chromosome 1 open reading frame 28	NM_024529	1.04
protein tyrosine phosphatase, non-receptor type 21	BF116199	1.04
discs, large (Drosophila) homolog 1	AW139131	1.04
non-metastatic cells 3, protein expressed in	AL031718	1.04
latent transforming growth factor beta binding protein 1	NM_006827	1.04
ATP synthase, H+ transporting, mitochondrial F1 complex, alpha subunit, isoform 1, cardiac muscle	AU140931	1.04
hemochromatosis	AF144244	1.04
dimethylarginine dimethylaminohydrolase 2	NM_013974	1.04
Consensus includes gb:AI692428 /FEA=EST /DB_XREF=gi:4969768 /DB_CLONE=IMAGE:2336854 /UG=Hs.184019 Homo sapiens clone 23551	AI692428	1.04
bile acid Coenzyme A: amino acid N-acyltransferase (glycine N-choloyltransferase)	NM_001701	1.04
DIPB protein	BF431488	1.04
serum/glucocorticoid regulated kinase	NM_005827	1.04
BET1 homolog (S. cerevisiae)	BC008899	1.04
TAF9-like RNA polymerase II, TATA box binding protein (TBP)-associated factor, 31 kD	AF077053	1.04
immunoglobulin kappa constant	X72475	1.04
RAO31-interacting protein	BE966146	1.04
ectonucleoside triphosphate diphosphohydrolase 1	U87967	1.04
retinal outer segment membrane protein 1	NM_000327	1.04
integrator receptor matrix proteoglycan 200	NM_016247	1.04
hypothetical protein FLJ21919	NM_023015	1.04
polyglutamine binding protein 1	AB041834	1.04
hypothetical protein FLJ10199	NM_018022	1.04
myeloid leukemia factor 2	NM_005439	1.04

gamma-tubulin complex protein 2	BF002130	1.04
glycine receptor, beta	AF094754	1.04
excision repair cross-complementing rodent repair deficiency, complementation group 4	U64315	1.04
phorbol-like protein MOS019	BFS08948	1.04
melanoma antigen, family A, 12	U82671	1.04
hypothetical protein FLJ10483	NM_018108	1.04
KIAA0795 protein	NM_025010	1.04
plaxin B2	BC004542	1.04
KIAA0295 protein	AB002293	1.04
Consensus includes gb:AA721025 /FEA=EST /DB_XREF=gi:2737160 /DB_XREF=est:ns8905.s1 /CLONE=IMAGE:1269464 /UG=Hs.293253 ESTs	AA721025	1.04
hypothetical protein MY014	BF938727	1.04
tumor necrosis factor, alpha-induced protein 6	AW188198	1.04
KIAA0633 gene product	NM_014645	1.04
PR domain containing 2, with ZNF domain	U23736	1.04
gb:M26463.1 /DEF=Homo sapiens immunoglobulin mu chain antibody MO30 (IgM) mRNA, complete cds. /FEA=CDS /GEN=IgM /PROD=immunoglobulin mu chain antibody	M28463	1.04
prostate epithelium-specific Ets transcription factor	A1307915	1.04
synaptophysin 1	BC000731	1.04
SUMO-1-specific protease	NM_015571	1.04
FYN binding protein (FYB-120/130)	A633888	1.04
hypothetical protein FLJ10901	NM_018265	1.04
cadherin 3, type 1, P-cadherin (placental)	NM_001793	1.04
glutaredoxin (thioltransferase)	NM_002084	1.04
lactoferrin (lactoferrin)	BF665706	1.04
zinc finger protein 219	NM_016423	1.04
calcium channel, voltage-dependent, alpha 1G subunit	AF126966	1.04
heparan sulfate proteoglycan 2 (perlecan)	A1981033	1.04
Consensus includes gb:AK022309.1 /DEF=Homo sapiens cDNA FLJ12247 fls, clone MAMMA1001397. /FEA=mRNA /DB_XREF=gi:10433678 /UG=Hs.287506 Homo sapiens	AK022309	1.04
KIAA1888 protein	AW444969	1.04
hypothetical protein FLJ12618	NM_024884	1.03
microtubule-associated protein, RPIEB family, member 3	A185178	1.03
nuclear receptor subfamily 2, group C, member 2	NM_003288	1.03
CD209 antigen-like	NM_014257	1.03
Consensus includes gb:AU146794 /FEA=EST /DB_XREF=gi:11008315 /DB_XREF=est:AU146794 /CLONE=HEMBB1001536 /UG=Hs.287475 Homo sapiens cDNA FLJ121	AU146794	1.03
prolactin-induced protein	NM_002652	1.03
procollagen C-endopeptidase enhancer 2	NM_013363	1.03
hypothetical protein FLJ22569	NM_023925	1.03
soluble carrier family 24 (sodium/potassium/calcium exchanger), member 2	NM_020344	1.03
uncharacterized hypothetical protein HCDASE	NM_018479	1.03
CGI-96 protein	AL589666	1.03
H2A histone family, member D	BF061074	1.03
homoeo box A9	U41813	1.03
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	NM_002810	1.03
Consensus includes gb:AL121994 /DEF=Human DNA sequence from clone RP4-781L3 on chromosome 1p34.3-36.11 Contains a pseudogene similar to IFITM3 (interferon	AL121994	1.03
Consensus includes gb:BF222018 /FEA=EST /DB_XREF=gi:1129195 /DB_XREF=est:7p41a04.x1 /CLONE=IMAGE:3648271 /UG=Hs.289803 ESTs. Moderately similar to BF222018	BF222018	1.03
microseminoprotein, beta-	NM_002443	1.03
KIAA1654 protein	AB051441	1.03
bone morphogenetic protein 2	NM_001200	1.03
collagen, type V, alpha 1	M76729	1.03
cytidine monophosphate-N-acetylneuraminic acid hydroxylase (CMP-N-acetylneuraminic acid hydroxylase)	AF074480	1.03
adipose differentiation-related protein	BC005127	1.03
Consensus includes gb:L43577 /FEA=EST /DB_XREF=gi:899066 /DB_XREF=est:L43577 /CLONE=EST500 /UG=Hs.82171 Homo sapiens (clone 115392) mRNA	L43577	1.03
B-cell linker	NM_013314	1.03
KIAA0284 protein	AB006622	1.03
uncharacterized hypothetical protein HT010	NM_018471	1.03
solute carrier family 6 (neurotransmitter transporter, serotonin), member 4	NM_001045	1.03
eosinophil peroxidase	X14346	1.03
Consensus includes gb:A871747 /FEA=EST /DB_XREF=gi:4851478 /DB_XREF=est:wa0506.x1 /CLONE=IMAGE:2297218 /UG=Hs.81360 Homo sapiens cDNA: FLJ2195	A871747	1.03
chromosome 21 open reading frame 80	BC006626	1.03
bulky like protein 3	A1591305	1.03
hypothetical protein FLJ20972	NM_025030	1.03
glucosyltransferase 6 (alpha (1.3) glucosyltransferase)	M98625	1.03
Conserved gene telomeric to alpha globin cluster	BE791629	1.03
CGI-01 protein	AL048669	1.03

protein tyrosine phosphatase, receptor type, J	O37781	1.03
peptide YY	D13902	1.03
catenin (cadherin-associated protein), delta 1	NM_001331	1.03
mitochondrial intermediate peptidase	U80034	1.03
TAF4B RNA polymerase II, TATA box binding protein (TBP)-associated factor, 105 kD	U09321	1.03
laminin, alpha 4	U77706	1.03
histatin 3	NM_000200	1.03
receptor tyrosine kinase-like orphan receptor 1	NM_005012	1.03
fibulin 5	NM_006329	1.03
hypothetical gene MGC9753	BF033007	1.03
mannosidase, beta A, lysosomal	NM_005908	1.03
KIAA0877 protein	BF002844	1.03
Consensus includes gb:AW161628 /FEA=EST /DB_XREF=gi:6300659 /DB_XREF=est:au68b11.x1 /CLONE=IMAGE:2781405 /UG=Hs.21739 Homo sapiens mRNA; cDNA	AW161628	1.03
KIAA1354 protein	AW138584	1.03
bone morphogenetic protein 7 (osteogenic protein 1)	AL157414	1.03
chromosome 1 open reading frame 24	NM_022083	1.03
chemokine (C-C motif) receptor 2	NM_000647	1.03
agouti related protein homolog (mouse)	NM_001138	1.03
retinol binding protein, 3, interstitial	J03912	1.03
nuclear protein ANKT	NM_016359	1.03
Kruppel-like factor 2 (lung)	NM_016270	1.03
KIAA1115 protein	BC002789	1.03
caveolin 1, caveolae protein, 22kD	AU147399	1.03
phospholipase D1, phosphatidylcholine-specific	AJ276230	1.03
ATPase, Class V, type 10D	AU78147	1.03
adrenergic, beta-2-, receptor, surface	NM_000024	1.03
hypothetical protein MGC2487	NM_023932	1.03
Consensus includes gb:AU155105 /FEA=EST /DB_XREF=gi:11016626 /DB_XREF=est:AU155105 /CLONE=OVARC:1001000 /UG=Hs.301348 Homo sapiens cDNA FLJ13;	AU155105	1.03
phospholipase A2, group IVB (cytosolic)	N71116	1.03
wingless-type MMTV integration site family, member 2B	AB045117	1.03
myo48 protein	AF063608	1.03
neuronal cell adhesion molecule	U55258	1.03
ubiquitin-conjugating enzyme E2D 2 (UBC4/S homolog, yeast)	BF196842	1.03
CLIP-associated protein 2	AL137636	1.03
RAS p21 protein activator (GTPase activating protein) 3 (ins1(1.3.4.5)/P4-binding protein)	NM_007368	1.03
Consensus includes gb:AK00144.1 /DEF=Homo sapiens cDNA FLJ20137 fls, clone COL07137. /FEA=mRNA /DB_XREF=gi:7020040 /UG=Hs.274449 Homo sapiens cDN	AK00144	1.03
solute carrier family 25 (mitochondrial oxalocarboxylate carrier), member 21	NM_030631	1.03
NGF1-A binding protein 1 (EGF1 binding protein 1)	AF045452	1.03
ribothymid II	AL514285	1.03
thyrotropin-releasing hormone receptor	O16845	1.03
PDZ-LIM protein mystique	NM_021630	1.03
centrosomal P4, 1-associated protein	NM_018451	1.03
UDP-N-acetyl-alpha-D-galactosamine polypeptide N-acetylgalactosaminyltransferase 6 (GalNAc-T6)	NM_007210	1.03
major vault protein	NM_017458	1.03
Sp1 transcription factor	AU121035	1.03
diaphorase (NADH) (cytochrome b-5 reductase)	NM_000398	1.03
signal recognition particle receptor (docking protein)	BC474341	1.03
nuclear factor (erythroid-derived 2)-like 3	NM_004289	1.03
Consensus includes gb:AW071997 /FEA=EST /DB_XREF=gi:6026995 /DB_XREF=est:ws5709.x1 /CLONE=IMAGE:2501321 /UG=Hs.328248 Homo sapiens mRNA; cDNA	AW071997	1.03
Consensus includes gb:AF216683 /DEF=Homo sapiens Interleukin-1 receptor antagonist homolog 1 (IL1HY1) gene, complete cds /FEA=mRNA, 1 /DB_XREF=gi:8572054 /	AF216683	1.03
Cluster ind. H59033/y40e10.r1 Homo sapiens cDNA, 5' end /clone=IMAGE:207786 /clone_and=5' /gb=H59033 /gi=1011865 /ug=Hs.6700 /len=604	H59033	1.03
elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1	H93026	1.03
deafness locus associated putative guanine nucleotide exchange factor	NM_012139	1.03
cell adhesion molecule with homology to L1CAM (close homologue of L1)	NM_006614	1.03
highly expressed in cancer, rich in leucine heptad repeats	NM_006101	1.03
hypothetical protein FLJ11730	NM_022736	1.03
hypothetical protein MGC4342	NM_024329	1.03
hypothetical protein FLJ13189	NM_024682	1.03
KIAA0564 protein	A1744628	1.03
peroxisomal long-chain acyl-coA thioesterase	NM_006821	1.03
adaptor-related protein complex 3, mu 2 subunit	NM_006803	1.03
hypothetical protein MGC11242	NM_024320	1.03
Fc fragment of IgG, low affinity Iib, receptor for (CD32)	U80640	1.03

Langierhans cell specific c-type lectin
 insulin upstream factor 1
 KIAA1193 protein
 Fc fragment of IgA receptor for
 underglycosylated hypobalaminus protein HT013
 hypothetical protein FLJ20605
 phosphatidic acid phosphatase type 2C
 odorant marker protein
 hypothetical protein MGC5590
 NTT5 protein
 gb:BC000110.1 /DEF=Homo sapiens. Similar to E2F transcription factor 5, clone MGC:1262, mRNA, complete cds. /FEA=mRNA /PROD=Similar to E2F transcription factor
 latent transforming growth factor beta binding protein 4
 hypothetical protein FLJ11736
 LOC86523
 tenomodulin protein
 hepatocyte growth factor (hepatopoietin A; scatter factor)
 KIAA1487 protein
 RAD51-like 3 (S. cerevisiae)
 crumba homolog 1 (Drosophila)
 chromosome 11 open reading frame 21
 ring finger protein 34
 hypothetical protein FLJ14753
 PTPRF interacting protein, binding protein 1 (liprin beta 1)
 DKFZP586F1918 protein
 deiodinase, liothyronine, type II
 KIAA0624 protein
 presenilin 2 (Alzheimer disease 4)
 KIAA0963 protein
 hypothetical protein PRQ2792
 KIAA0304 gene product
 chloride channel Kb
 peptidoglycan recognition protein
 growth factor receptor-bound protein 10
 Consensus includes gb:A1700633 /FEA=EST /DB_XREF=gi:4988533 /DB_XREF=gi:4988533 /DB_XREF=est:wa39g03.x1 /CLONE=IMAGE:2343412 /UG=Hs.286232 Homo sapiens cDNA: FLJ221
 solute carrier family 28 (sodium-coupled nucleoside transporter), member 3
 ring finger protein 24
 hypothetical protein FLJ21031
 translocating chain-associating membrane protein
 receptor (calicrin) activity modifying protein 2
 CD81 antigen (target of antiproliferative antibody 1)
 zinc finger protein 220
 cytosolic ovarian carcinoma antigen 1
 excision repair cross-complementing rodent repair deficiency, complementation group 2 (xeroderma pigmentosum D)
 heparan sulfate (glucosamine) 3-O-sulfotransferase 3A1
 metal-regulatory transcription factor 1
 DiGeorge syndrome gene H
 methyl-CpG binding domain protein 4
 hypothetical protein FLJ13465
 DKFZP586F1918 protein
 thyroid hormone receptor-associated protein, 95-kD subunit
 bromodomain-containing 1
 Consensus includes gb:BG500611 /FEA=EST /DB_XREF=gi:13462128 /DB_XREF=est:602546082F1 /CLONE=IMAGE:4668261 /UG=Hs.82719 Homo sapiens mRNA; cdi
 retin
 phosphoinositide-3-kinase, catalytic, delta polypeptide
 ATP-binding cassette, sub-family C (CFTR/MRP), member 9
 Consensus includes gb:AK023890.1 /DEF=Homo sapiens cDNA FLJ13628 fis, clone PLACE1011054. /FEA=mRNA /DB_XREF=gi:10435692 /UG=Hs.306656 Homo sapiens
 COX11 homolog, cytochrome c oxidase assembly protein (yeast)
 MAP-kinase activating death domain
 blepharophimosis, epicanthus inversus and ptosis, candidate 1
 hypothetical protein PRO2834
 Fc fragment of IgG, low affinity IIb, receptor for (CD16)
 hypothetical protein
 ATP-binding cassette, sub-family E (OABP), member 1

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ATP-dependent RNA helicase	AF077033	1.02
synaptonemal complex protein 2	NM_014258	1.02
LDL receptor adaptor protein	AL545035	1.02
KIAA0586 gene product	NM_014749	1.02
KIAA0700 protein	AW450345	1.02
Consensus includes gb:BG105365 /FEA=EST /DB_XREF=est:602312473F1 /CLONE=IMAGE:4422043 /UG=Hs.23348 S-phase kinase-associated BG105365	AK023365	1.02
Consensus includes gb:AK023365.1 /DEF=Homo sapiens cDNA FLJ13303 f1s, clone OVARC1001372, highly similar to Homo sapiens liprin-alpha4 mRNA. /FEA=mRNA /C	AK023365	1.02
Meiosis associated gene	BF342851	1.02
hypothetical protein FLJ23138	AW000928	1.02
tetratricopeptide repeat domain 3	AW510696	1.02
hypothetical protein MGC3282	NM_024029	1.02
erythroblast membrane-associated protein	NM_018538	1.02
ubiquitin-conjugating enzyme E2D 1 (UBC4/5 homolog, yeast)	BC005980	1.02
K14-type splicing regulatory protein (FUSE binding protein 2)	AI933301	1.02
paired box gene 3 (Waaerdenburg syndrome 1)	NM_013942	1.02
KIAA0532 protein	AI052003	1.02
sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4G	NM_017893	1.02
purinergic receptor P2X, ligand-gated ion channel, 3	NM_002559	1.02
Bardet-Biedl syndrome 4	AI813772	1.02
KIAA0401 protein	AB007861	1.02
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson	NM_022975	1.02
DKFZp566D133 protein	BC436400	1.02
hypothetical protein FLJ20871 similar to FSP27	NM_022094	1.02
neurogenic differentiation 6	NM_022728	1.02
pregnancy specific beta-1-glycoprotein 3	BC005824	1.02
phosphatidylinositol binding clathrin assembly protein	AL135735	1.02
solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2	AK5105	1.02
zinc finger protein 48 (ZFP)	NM_008977	1.02
Consensus includes gb:AF052119.1 /DEF=Homo sapiens clone 23622 mRNA sequence /FEA=mRNA /DB_XREF=gi:3360428 /UG=Hs.151608 Homo sapiens clone 23622 AF052119	AF052119	1.02
purinergic receptor P2X-like 1, orphan receptor	NM_005446	1.02
VIR-like 1	NM_020633	1.02
KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2	NM_068854	1.02
fibroblast growth factor receptor 2 (bacteria-expressed kinase, keratinocyte growth factor receptor, craniofacial dysostosis 1, Crouzon syndrome, Pfeiffer syndrome, Jackson	M87771	1.02
5' nucleoidase (CD73)	NM_002526	1.02
DNA (cytosine-5)-methyltransferase 2	AJ223333	1.02
bagpope homeobox homolog 1 (Drosophila)	NM_001189	1.02
Consensus includes gb:AK022254.1 /DEF=Homo sapiens cDNA FLJ12192 f1s, clone MAMMA1000851. /FEA=mRNA /DB_XREF=gi:10433612 /UG=Hs.288451 Homo sapie	AK022254	1.02
sialyltransferase 4C (beta-galactosidase alpha-2,3-sialyltransferase)	AK021929	1.02
aspartate beta-hydroxylase	NM_020164	1.02
Consensus includes gb:AL049387.1 /DEF=Homo sapiens mRNA; cDNA DKFZp586N1918 (from clone DKFZp586N1918) /FEA=mRNA /DB_XREF=gi:4500178 /UG=Hs.16	AL049387	1.02
LIM domain only 7	NM_005358	1.02
Consensus includes gb:BF431965 /FEA=EST /DB_XREF=gi:11444079 /DB_XREF=est:na83409.x1 /CLONE=IMAGE:3274409 /UG=Hs.153106 Homo sapiens clone 2372 BF431965	BF431965	1.02
casepase 10, apoptosis-related cysteine protease	AF111345	1.02
carbohydrate (N-acetylglucosamine 4-0) sulfotransferase 6	NM_022467	1.02
ten-1 homolog b (C. elegans)	NM_015322	1.02
glucose-6-phosphatase, Transport (glucose-6-phosphate) protein 1	NM_001467	1.02
zinc finger, imprinted 2	NM_015383	1.02
catenin (cadherin-associated protein), delta 1	AB023382	1.02
phosphoinositide-3-kinase, regulatory subunit, polypeptide p101	NM_014308	1.02
retinoblastoma binding protein 5	NM_005057	1.02
bone morphogenetic protein 8 (osteogenic protein 2)	NM_001720	1.02
dermatopontin	NM_001937	1.02
antyrin-like with transmembrane domains 1	NM_007332	1.02
glucose-6-phosphatase, catalytic (glycogen storage disease type I, von Gierke disease)	NM_000151	1.02
retinitis pigmentosa 2 (X-linked recessive)	NM_006915	1.02
lymphoid nuclear protein related to Ar-4	AI90465	1.02
tumor necrosis factor receptor superfamily, member 5	BF664114	1.02
ring finger protein 13	NM_007282	1.02
hypothetical protein FLJ20421	NM_017813	1.02
ATPase, Na+/K+ transporting, alpha 2 (*) polypeptide	NM_024719	1.02
protein phosphatase 3 (formerly 2B), regulatory subunit B (19kD), alpha isoform (calcieneurin B, type I)	NM_000702	1.02
farnesyltransferase, CAAX box, beta	AL544951	1.02
	NM_002028	1.02

KIAA0781 protein
protein O-ucosyltransferase 1
Cluster incl. AL120741:DKFZp782K231_r1 Homo sapiens cDNA, 5' end /clone=DKFZp782K231 /clone_end=5 /gb=AL120741 /gi=5926641 /ug=Hs.8859 /len=707
aa homologous factor
neutensin
Consensus includes gb:AC005614 /DEF=Homo sapiens chromosome 19, cosmid F23269 /FEA=CDS /DB_XREF=gi:3540175 /UG=Hs.287391 Homo sapiens chromosome AC005614
Consensus includes gb:AF090100.1 /DEF=Homo sapiens clone IMAGE 23915 /FEA=mrna /DB_XREF=gi:4063635 /UG=Hs.170241 Homo sapiens clone IMAGE 23915
KIAA1097 protein
chromosome 11 open reading frame 17
hypothetical protein
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 13
p21/Cdc42/Rac1-activated kinase 1 (STE20 homolog, yeast)
SH3-domain binding protein 2
gb:NM_023939.1 /DEF=Homo sapiens hypothetical protein MGC2752 (MGC2752), mRNA, /FEA=mrna /GEN=MGC2752 /PROD=hypothetical protein MGC2752 /DB_XREF=NM_023939
tumor necrosis factor (ligand) superfamily, member 15
Consensus includes gb:AI652058 /FEA=EST /DB_XREF=gi:4736037 /DB_XREF=est:w627411.x1 /CLONE=IMAGE 2306901 /UG=Hs.129061 Human Chromosome 16 BAC AI652058
KIAA0164 gene product
histone deacetylase 2
ret finger protein
glycerol kinase
KIAA0194 protein
PPAR binding protein
killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 3
DKFZP566E104 protein
interactin 1 (SH3 domain protein)
SPTF-associated factor 65 gamma
RAR-related orphan receptor A
ubiquitin specific protease 10
Consensus includes gb:AF035281.1 /DEF=Homo sapiens clone 23903 mRNA sequence, /FEA=mrna /DB_XREF=gi:2661032 /UG=Hs.12526 Homo sapiens clone 23903
SELENOPHOSPHATE SYNTHETASE 1
selenium binding protein 1
hypothetical protein FLJ20897
myeloid/lymphoid or mixed-lineage leukemia (trithorax homolog, Drosophila); translocated to, 3
hypothetical protein FLJ10890
pregnancy specific beta-1-glycoprotein 2
chromosome 12 open reading frame 3
ferritin, light polypeptide
ATP-binding cassette, sub-family C (CFTR/MRP), member 6
phosphatidylinositol transfer protein, membrane-associated
high-glucose-regulated protein 8
growth associated protein 43
sash-repeat protein
HIV TAT specific factor 1
purine-rich element binding protein A
KIAA0359 protein
CREB binding protein (Rubinstein-Taybi syndrome)
plasminogen activator, urokinase receptor
neutral sphingomyelinase
tumor necrosis factor receptor superfamily, member 6b, decoy
4-aminobutyrate aminotransferase
WD repeat domain 10
adenylate cyclase 3
lanosterol synthase (2,3-oxidosqualene-lanosterol cyclase)
connexin 59
hypothetical protein FLJ14360
platelet-derived growth factor receptor, alpha polypeptide
ncam
glutathione peroxidase 5 (epididymal androgen-related protein)
Snf2-related CBP activator protein
glycogenin 2
hypothetical protein FLJ13615
I-box and leucine-rich repeat protein 11
stromal antigen 1

AB018324 1.02
BC000582 1.02
AL120741 1.02
NM_012153 1.02
NM_006183 1.02
AC005614 1.02
AF090100 1.02
AK02864 1.02
NM_020642 1.02
NM_020313 1.02
AJ001698 1.02
U51120 1.02
AB000463 1.02
NM_023939 1.02
NM_065118 1.02
NM_014739 1.02
NM_001527 1.02
AL523814 1.02
X68285 1.02
D83778 1.02
AF055994 1.02
AC006293 1.02
AL110238 1.02
AF064243 1.02
AU154740 1.02
L14611 1.02
BG390445 1.02
AF035281 1.02
AV682679 1.02
NM_003944 1.02
BF690020 1.02
NM_004529 1.02
NM_018259 1.02
NM_031246 1.02
NM_020373 1.02
BF312331 1.02
NM_001171 1.02
NM_004910 1.02
NM_016258 1.02
NM_002045 1.02
NM_014467 1.02
NM_014500 1.02
BF739943 1.02
NM_015113 1.02
NM_004380 1.02
U06839 1.02
NM_016317 1.02
NM_016434 1.02
NM_000663 1.02
NM_018262 1.02
AF033861 1.02
D63807 1.02
NM_030772 1.02
N36751 1.02
NM_006208 1.02
NM_016637 1.02
NM_003996 1.02
AK023808 1.02
U94363 1.02
AF317887 1.02
BE675843 1.02
AW169948 1.02

acyl-Coenzyme A oxidase 1, palmitoyl
hypothetical protein FLJ20130
Consensus includes gb:AA534504 /FEA=EST /DB_XREF=gi:22718757 /DB_XREF=IMAGE-926246 /UG=Hs.8982 ESTs, Highly similar to KIAA1395
Janus kinase 1 (a protein tyrosine kinase)
cytochrome b-561
solute carrier family 24 (sodium/potassium/calcium exchanger), member 1
FAST kinase
cyclin-dependent kinase inhibitor 1C (p57, Kip2)
interleukin 15
kinase suppressor of ras
heparanase
KIAA0653 protein, type IV, alpha 1
Consensus includes gb:AI867981 /FEA=EST /DB_XREF=gi:5764799 /DB_XREF=est:wu1105.x1 /CLONE=IMAGE:2516673 /UG=Hs.115065 ESTs
chromosome 22 open reading frame 5
solute carrier family 4, sodium bicarbonate cotransporter, member 7
hypothetical protein FLJ11467
Consensus includes gb:AA769006 /FEA=EST /DB_XREF=gi:2820244 /DB_XREF=IMAGE:1318578 /UG=Hs.290921 ESTs, Highly similar to G027
heparan sulfate proteoglycan 2 (perlecan)
peptidyl-prolyl isomerase G (cyclophilin G)
cholecystokinin
msh homeo box homolog 2 (Drosophila)
Consensus includes gb:AK025422.1 /DEF=Homo sapiens cDNA: FLJ21769 fls, clone COL F7354, /FEA=mRNA /DB_XREF=gi:10437930 /UG=Hs.306810 Homo sapiens cDNA: AK025422
hypothetical protein FLJ12871
pleomorphic adenoma gene-like 2
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 4
Janus kinase 3 (a protein tyrosine kinase, leukocyte)
protein tyrosine phosphatase, non-receptor type 7
hypothetical protein FLJ22527
erythropoietin receptor
interferon (alpha, beta and omega) receptor 2
Consensus includes gb:AF070538.1 /DEF=Homo sapiens clone 24420 mRNA sequence, /FEA=mRNA /DB_XREF=gi:3387895 /UG=Hs.159408 Homo sapiens clone 24420 mRNA sequence: AF070538
alpha2,3-sialyltransferase
SH2 domain protein 1A, Duncan's disease (lymphoproliferative syndrome)
hyperpolarization activated cyclic nucleotide-gated potassium channel 2
Consensus includes gb:AK026682.1 /DEF=Homo sapiens cDNA: FLJ23029 fls, clone LING01883, /FEA=mRNA /DB_XREF=gi:10439598 /UG=Hs.308864 Homo sapiens cDNA: AK026682
hypothetical gene DKFZp434A1010
endothelial differentiation, sphingolipid G-protein-coupled receptor, 5
lysophospholipase 1
chromosome 21 open reading frame 8
serine (or cysteine) proteinase inhibitor, clade A (alpha-1 antitrypsin, antitrypsin), member 1
Consensus includes gb:AV647366 /FEA=EST /DB_XREF=gi:9868380 /DB_XREF=est:AV647366 /CLONE=GLCAXE09 /UG=Hs.282365 ESTs
solute carrier family 30 (zinc transporter), member 3
adenylyl cyclase-associated protein
Consensus includes gb:U82303.1 /DEF=Homo sapiens unknown protein mRNA, partial cds, /FEA=mRNA /PROD=unknown /DB_XREF=gi:1938329 /UG=Hs.123080 Homo sapiens clone 1938329 mRNA: U82303
presenilin 2 (Alzheimer disease 4)
phytanoyl-CoA hydroxylase (Refsum disease)
hypothetical protein from EUROIMAGE 2021883
Cluster Inc. AI198543:q49f11.x1 Homo sapiens cDNA, 3' end /clone=IMAGE-1753389 /clone_end=3 /gb=AI198543 /gi=3751149 /ug=Hs.8982 /len=794
alpha-actinin-2-associated LIM protein
DNA binding protein for surfactant protein B
KIAA1085 protein
discs, large (Drosophila) homolog 1
trypartite motif-containing 9
myosin VI
killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 4
tumor necrosis factor receptor superfamily, member 6b, decoy
interferon-stimulated protein, 15 kDa
KIAA0507 protein
transcobalamin I (vitamin B12 binding protein, R binder family)
pleomorphic adenoma gene-like 1
spem associated antigen 11
neurofibromin 2 (bilateral acoustic neuroma)

Consensus includes gb:AU147599 /FEA=EST /DB_XREF=gi:1009120 /DB_XREF=est:AU147599 /CLONE=MAMMA1001154 /UG=Hs.287501 Homo sapiens cDNA FLJ12
 galactose-4-epimerase, UDP. NM_000403 1.02
 tumor necrosis factor receptor superfamily, member 10b AF153887 1.02
 zinc finger protein NM_018651 1.02
 neuropathy target esterase NM_006702 1.02
 solute carrier family 16 (monocarboxylic acid transporters), member 3 AA527578 1.02
 myelin transcription factor 1 AB020842 1.02
 chymotrypsin-like AD72212 1.02
 ubiquitin specific protease 3 AF077040 1.02
 keratin, hair, basic, 1 NM_002281 1.02
 Sec23-interacting protein p125 AK001135 1.02
 interleukin 5 receptor, alpha M75914 1.02
 retinoic acid receptor responder (tazarotene induced) 1 NM_002888 1.02
 transmembrane 4 superfamily member 5 NM_003963 1.02
 nuclear LIM domain-interacting factor NM_021198 1.02
 ETAA16 protein NM_019002 1.02
 Consensus includes gb:AI992251 /FEA=EST /DB_XREF=gi:5839156 /DB_XREF=est:ws41d12.x1 /CLONE=IMAGE:2499767 /UG=Hs.184591 ESTs
 glycosylphosphatidylinositol specific phospholipase D1 AI992251 1.02
 Consensus includes gb:BF448703 /FEA=EST /DB_XREF=gi:11514872 /DB_XREF=est:7n93902.x1 /CLONE=IMAGE:3572019 /UG=Hs.59728 Homo sapiens mRNA; cDNA BF448703 1.02
 crystallin, beta A1 NM_005208 1.02
 KIAA0763 gene product AI652645 1.02
 mucin 3A, intestinal AF007194 1.02
 Consensus includes gb:AA595276 /FEA=EST /DB_XREF=gi:2410626 /DB_XREF=est:tno35a05.s1 /CLONE=IMAGE:1102640 /UG=Hs.243804 Homo sapiens cDNA FLJ136
 v-src sarcoma (Schmidt-Ruppin A-2) viral oncogene homolog (avian) AK024281 1.02
 RAB3B, member RAS oncogene family NM_002867 1.02
 natural killer-tumor recognition sequence AI361805 1.02
 GATA binding protein 4 AV700724 1.02
 DKFZP434J154 protein AK024279 1.02
 MAD, mothers against decapentaplegic homolog 6 (Drosophila) NM_005585 1.02
 growth differentiation factor 3 NM_020634 1.02
 ribosomal protein S12 BC002550 1.02
 AT2 receptor-interacting protein 1 AI695017 1.02
 KIAA0692 protein AI924382 1.02
 hypothetical protein DKFZp564B1162 NM_031305 1.02
 leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 7 AF212842 1.02
 Consensus includes gb:AW450910 /FEA=EST /DB_XREF=gi:6991868 /DB_XREF=est:UH-H-B10-alle-01-Q-UI.s1 /CLONE=IMAGE:2737225 /UG=Hs.74101 spleen tyrosine
 gb:LO8961.1 /DEF=Homo sapiens transmembrane tyrosine kinase mRNA, complete cds. /FEA=CDS /PROD=tyrosine kinase /DB_XREF=gi:697614 /FL-gb:LO8961.1
 neuro-oncological ventral antigen 2 L08961 1.02
 Consensus includes gb:R48779 /FEA=EST /DB_XREF=gi:810805 /DB_XREF=est:ty69c07.s1 /CLONE=IMAGE:153996 /UG=Hs.293845 ESTs, Weakly similar to B32688 bt
 candidate tumor suppressor p33 ING1 homolog NM_002516 1.02
 RYK receptor-like tyrosine kinase AA887063 1.02
 runt-related transcription factor 1 (acute myeloid leukemia 1; aml1 oncogene) BG032035 1.02
 sialophorin (gp115, leukostatin, CD43) AF312387 1.02
 PDZ domain containing 1 J04168 1.02
 capping protein (actin filament) muscle Z-line, beta NM_002614 1.02
 serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 6 U03271 1.02
 tumor necrosis factor receptor superfamily, member 12 (translocating chain-association membrane protein) BC004948 1.02
 microtubule-associated protein, RP/EB family, member 2 BE671156 1.02
 claudin 3 NM_001306 1.02
 renin binding protein NM_002910 1.02
 HIV-1 Rev binding protein A89512 1.02
 natriuretic receptor 2 (cardiac) NM_001035 1.02
 Consensus includes gb:BF939092 /FEA=EST /DB_XREF=gi:12356412 /DB_XREF=est:705c01.x1 /CLONE=IMAGE:3706753 /UG=Hs.136090 ESTs
 alanine-glyoxylate aminotransferase (oxalosis I; hyperoxaluria I; glycolicaciduria; serine-pyruvate aminotransferase) BF939092 1.02
 prostaglandin E receptor 1 (subtype EP1), 42x0 D13368 1.02
 calcium binding protein NM_000655 1.02
 hypothetical protein PRO0756 NM_018583 1.02
 hypothetical protein NM_018855 1.02
 old astrocyte specifically induced substance Z62180 1.02
 semaphorin 5A AF055009 1.02
 aldolase B, fructose-bisphosphate NM_003966 1.02
 LIM domain protein BF195998 1.02
 apoptosis inhibitor 5A NM_003687 1.02
 AF229254 1.02

ribosomal protein L31
 hypothetical protein FLJ13171
 solute carrier family 21 (organic anion transporter), member 9
 cytochrome P450, subfamily XXVIA, polypeptide 1
 protein phosphatase 1, regulatory (inhibitor) subunit 8
 Consensus includes gb:AL049351.1 /DEF=Homo sapiens mRNA: cDNA DKFZp566C093 (from clone DKFZp566C093). /FEA=mRNA /DB_XREF=gi:4500138 /UG=Hs.3020
 erythrocyte protein band 4.1-like 4
 Golgi apparatus protein 1
 small nuclear RNA activating complex, polypeptide 5, 19KD
 DKFZP568M1523 protein
 phosphoinositide-3-kinase, catalytic, gamma polypeptide
 nuclear mitotic apparatus protein 1
 IGF-II mRNA-binding protein 3
 poliovirus receptor
 KIAA0753 gene product
 hemochromatosis
 hypothetical protein
 likely ortholog of mouse ubiquitin-conjugating enzyme E2-230K
 Consensus includes gb:BG420747 /FEA=EST /DB_XREF=gi:13327253 /DB_XREF=est:602448537 /1 /CLONE=IMAGE:4586984 /UG=Hs.259677 ESTs, Weakly similar to : BG420747
 anti-Müllerian hormone receptor, type II
 AND-1 protein
 palmitoyl-protein thioesterase 2
 KIAA0719 protein
 engulfment and cell motility 2 (ced-12 homolog, C. elegans)
 centrin, EF-hand protein, 2
 cytochrome c-like antigen
 thromboxane A2 receptor
 cathrin, light polypeptide (Lcb)
 heterogeneous nuclear ribonucleoprotein D-like
 paraoxonase 3
 glypican 4
 methyl-CpG binding domain protein 2
 dyskeratosis congenita 1, dyskerin
 keratin 6B
 nuclear protein 3 (apoptosis repressor with CARD domain)
 glucosidase, beta: add (includes glucosylceramidase)
 chloride intracellular channel 5
 natural killer-tumor recognition sequence
 gollioth protein
 PAN binding protein 9
 tanatodictin (beta)-like 1
 Cluster Incl. A1744451.wf89d08.x1 Homo sapiens cDNA, 3' end /clones=IMAGE-2362787 /clone_end=3 /gb=A1744451 /gi=5112739 /ug=Hs.124820 /len=584
 protein predicted by clone 23733
 low density lipoprotein receptor-related protein 3
 microfilament-associated protein 3
 KIAA0630 protein
 fibroblast growth factor receptor 4
 RAB5A, member RAS oncogene family
 SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily c, member 2
 CREB-induced protein
 hypothetical protein FLJ12781
 adaptor-related protein complex 4, mu 1 subunit
 amyloid beta (A4) precursor protein-binding, family A, member 3 (X11-like 2)
 exosomes (multiple) 2
 putative dimethyladenosine transferase
 hypothetical protein FLJ20725
 Consensus includes gb:AF052100.1 /DEF=Homo sapiens clone 23645 mRNA sequence /FEA=mRNA /DB_XREF=gi:3360407 /UG=Hs.6651 Homo sapiens clone 23645 m
 testis intracellular mediator protein
 Jun B proto-oncogene
 growth hormone 2
 serine/threonine kinase 13 (aurora/PL1-like)
 coagulation factor V (procoagelin, labile factor)
 TdT binding protein

F-box only protein 3
neuron-specific protein
hypothetical protein FLJ13204
transcription factor 12 (HTF4, helix-loop-helix transcription factors 4)
endoplasmic reticulum oxidoreductin 1-Lbeta
KIAA0038 gene product
cytochrome P450, subfamily XXVIII (steroid 27-hydroxylase, cerebrotendinous xanthomatosis), polypeptide 1
N-myristyltransferase 2
adrenomedullary alpha-2C₁ receptor
PAB38, member PAS oncogene family
hypothetical protein FLJ12401
P4508 binding protein 3 (25kD)
Consensus includes gb:AL031602 /DEF=Human DNA sequence from clone RP5-1174N9 on chromosome 1p34.1-35.3. Contains the gene for a novel protein with IBR dom
lysine 3-monoxygenase/hydrophobic 5-monoxygenase activation protein, theta polypeptide
KIT ligand
POU domain, class 3, transcription factor 4
Consensus includes gb:AU144900 /FEA=EST /DB_XREF=gi:11006421 /DB_XREF=est:AU144900 /CLONE=HEMBA1003327 /UG=Hs.274233 Homo sapiens cDNA FLJ10
T-cell leukemia/lymphoma 6
KIAA0774 protein
tRNA selenocysteine associated protein
growth factor receptor-bound protein 7
hypothetical protein MGC2716
Consensus includes gb:AA205593 /FEA=EST /DB_XREF=gi:1803801 /DB_XREF=est:zq6806.s1 /CLONE=IMAGE:646782 /UG=Hs.26837 Homo sapiens mRNA: cDNA DI
BE994361
melanoma cell adhesion molecule
hypothetical protein FLJ10408
wingless-type MMTV integration site family, member 5B
NIMA (never in mitosis gene a)-related kinase 1
hypothetical protein FLJ22794
fibulin 1
G protein-coupled receptor kinase 5
major histocompatibility complex, class II, DO alpha
KIAA1109 protein
olfactory receptor, family 1, subfamily F, member 1
mitogen-activated protein kinase kinase kinase 10
hypothetical protein MGC2550
toll-like receptor 3
KIAA0993 protein
CD5 antigen (p58-62)
MAD, mothers against decapentaplegic homolog (Drosophila) interacting protein, receptor activation anchor
actin-like 7B
hypothetical protein DKFZp586E1923
patched homolog (Drosophila)
Consensus includes gb:BE967207 /FEA=EST /DB_XREF=gi:11773627 /DB_XREF=est:601861094R1 /CLONE=IMAGE:3916174 /UG=Hs.165580 ribosomal protein S13
nuclear receptor subfamily 2, group F, member 6
thimet oligopeptidase 1
A kinase (PRKA) anchor protein (ytlaa) 9
CGI-48 protein
hypothetical protein MGC4845
cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 9
likely ortholog of preimplantation protein 3
methyl-CpG binding domain protein 2
lectin, mannose-binding, 1
Consensus includes gb:AI689947 /FEA=EST /DB_XREF=gi:4834721 /DB_XREF=est:wb68d12.x1 /CLONE=IMAGE:2312567 /UG=Hs.71969 Homo sapiens mRNA: cDNA [AI689947
protein phosphatase 1, regulatory (inhibitor) subunit 3A (glycogen and sarcoplasmic reticulum binding subunit, skeletal muscle)
RNA, U transporter 1
hypothetical protein FLJ10525
KIAA0783 gene product
KIAA0609 protein
inter-alpha (globulin) inhibitor H4 (plasmin-Kallikrein-sensitive glycoprotein)
gb:BC001279.1 /DEF=Homo sapiens. Similar to KIAA0626 gene product, clone MGC:5129. mRNA, complete cds. /FEA=mRNA /PROD=Similar to KIAA0626 gene product
transducin-like enhancer of split 2 (Elsp1) homolog (Drosophila)
Gem-interacting protein
ATP-binding cassette, sub-family C (CFTR/MRP), member 1

sec22 homolog
CD8 antigen, beta polypeptide 1 (p37)
5,10-methylenetetrahydrofolate synthetase (5-formyltetrahydrofolate cyclo-ligase)
basic leucine zipper transcription factor, ATF-like
major histocompatibility complex, class I-like sequence
lipophilin A (ulceroglobulin family member)
natural killer cell group 7 sequence
alpha-2-glycoprotein 1, zinc
nuclear phosphoprotein similar to S. cerevisiae PWP1
Rhesus blood group-associated glycoprotein
annexin A1
voltage-dependent anion channel 1 pseudogene
hypothetical protein MGC17330
hypothetical protein
thiamin pyrophosphokinase 1
integrin, beta 1 (fibronectin receptor, beta polypeptide, antigen CD29 includes MDF2, MSK12)
TAF_{II}25 RNA polymerase II, TATA box binding protein (TBP)-associated factor, 31 kD
MLL, septin-like fusion
Consensus includes gb:BG151527/FEA=EST/DB_XREF=gi:1263557/DB_XREF=est:nac28d07.x1/CLONE=IMAGE:3394597/UG=Hs.128095 ESTs, Moderately similar
solute carrier family 7 (cationic amino acid transporter, y⁺ system), member 6
guanine nucleotide binding protein (G protein), q polypeptide
Cip1-interacting zinc finger protein
fodillin 1
protein phosphatase 1, regulatory subunit 10
solute carrier family 7 (cationic amino acid transporter, y⁺ system), member 6
methylenetetrahydrofolate synthetase 1 (alpha)
ring finger protein 15
Consensus includes gb:A693140/FEA=EST/DB_XREF=gi:4970480/DB_XREF=est:wd68a05.x1/CLONE=IMAGE:2336720/UG=Hs.22907 Homo sapiens mRNA; cDNA
collator required for Sp1 transcriptional activation, subunit 9 (33kD)
neuro-oncological ventral antigen 1
GREB1 protein
alpha-methylacyl-CoA racemase
BUB3 budding uninhibited by benzimidazoles 3 homolog (yeast)
T. brachyury homolog (mouse)
hypothetical protein P15-2
hypothetical protein
neural proliferation, differentiation and control, 1
hypothetical protein FLJ10615
hypothetical protein DKFZp434G131
HCR (e-helix coiled-coil rod homologue)
penicillin-related gene, rapidly induced by IL-1 beta
KIAA0471 gene product
hypothetical protein DKFZp547D104
cellular repressor of E1A-stimulated genes
a disintegrin and metalloproteinase domain 15 (metagridin)
apical cord-derived growth factor-B
hypothetical protein FLJ11286
hypothetical protein FLJ20641
hypothetical protein FLJ12895
glial fibrillary acidic protein
bromodomain adjacent to zinc finger domain, 1B
ELL-RELATED RNA POLYMERASE II, ELONGATION FACTOR
G protein-coupled receptor 45
Consensus includes gb:AK025313.1 IDEF-Homo sapiens cDNA; FLJ21660 fis, clone COL08818./FEA=mRNA/DB_XREF=gi:10437803/UG=Hs.287678 Homo sapiens cDNA; AK025313
solute carrier family 2 (facilitated glucose transporter) member 8
interferon-stimulated transcription factor 3, gamma (48kD)
Janus kinase 3 (e protein tyrosine kinase, leukocyte)
hypothetical protein FLJ13725
zona pellucida binding protein
HSPC056 protein
hypothetical protein FLJ10211
preludin 4
Cluster Ind. A174083:wc38b04.x1 Homo sapiens cDNA, 3 end lclone=IMAGE:2317231/lclone_end=3 /gb=A1744083 /gi=5112371 /ug=Hs.65406 llen=608

myosin VA (heavy polypeptide 12, myosin)
 DEAD(H) (Asp-Glu-Ala-Asp/His) box polypeptide 38
 glucagon-like peptide 1 receptor
 endothelin receptor type B
 complement component 8, gamma polypeptide
 ubiquitination factor E4B (UFD2 homolog, yeast)
 stannin
 CGI-40 protein
 pyruvate dehydrogenase phosphatase
 hypothetical protein DKFZp547D1014
 Consensus includes gb:AL356602.1 /DEF=Homo sapiens mRNA; cDNA DKFZp547D146 (from clone DKFZp547D146) /FEA=mRNA /DB_XREF=gi:8655670 /UG=Hs.3065
 KIAA0545 protein
 glutathione S-transferase M1
 G protein-coupled receptor 68
 adenylate cyclase 1 (brain)
 leucine-rich repeat protein, neuronal 3
 hepy-RNA synthetase
 hypothetical protein FLJ20552
 Consensus includes gb:A1927701 /FEA=EST /DB_XREF=gi:5663665 /DB_XREF=est:wo65h11.x1 /CLONE=IMAGE:2462181 /UG=Hs.25307 Homo sapiens clone 24812 ml /A1927701
 LIM and cysteine-rich domains 1
 3-oxoacid CoA transferase
 Consensus includes gb:AW969913 /FEA=EST /DB_XREF=gi:8159737 /DB_XREF=est:EST381991 /UG=Hs.105121 ESTs
 calbindin 3, (vitamin D-dependent calcium binding protein)
 Consensus includes gb:A095896 /FEA=EST /DB_XREF=gi:3434872 /DB_XREF=est:qb21a10.x1 /CLONE=IMAGE:1696890 /UG=Hs.301463 Human Chromosome 16 BAC A095896
 N-ethylmaleimide-sensitive factor attachment protein, alpha
 chromosome 11 open reading frame 15
 synaptosomal-associated protein, 23kD
 ALM-1 protein
 hypothetical protein MGC14817
 Consensus includes gb:AF19846.1 /DEF=Homo sapiens PRO1474 mRNA, complete cds. /FEA=mRNA /PROD=PRO1474 /DB_XREF=gi:7770128 /UG=Hs.324925 LATS1 /AF19846
 DisGeorge syndrome critical region gene 6
 vascular Rab-GAP/TBC-containing
 KIAA0663 gene product
 colipase, pancreatic
 SMA5
 lysosomal-associated membrane protein 1
 propionyl Coenzyme A carboxylase, alpha polypeptide
 L-complex-associated-testis-expressed 1-like 1
 DKFZP434J214 protein
 myosin IB
 CASP8 and FADD-like apoptosis regulator
 Rab geranylgeranyltransferase, beta subunit
 hydroxysteroid (17-beta) dehydrogenase 4
 KIAA1088 protein
 ankyrin repeat and SOCS box-containing 1
 hypothetical protein MGC5178
 zinc finger protein 134 (clone pHZ-15)
 G protein-coupled receptor, family C, group 5, member D
 Consensus includes gb:X81695.1 /DEF=H.sapiens GENX-5624 mRNA, 3 UTR. /FEA=mRNA /DB_XREF=gi:2370158 /UG=Hs.194765 H.sapiens GENX-5624 mRNA, 3 UTR /X81695
 DKFZP434J082 protein
 aspartate carboxylase (aminocyclase 2, Carnahan disease)
 succinate dehydrogenase complex, subunit B, iron sulfur (lp)
 MAX interacting protein 1
 peroxisomal biogenesis factor 6
 chromosome 17 open reading frame 1A
 siruin silent mating type information regulation 2 homolog 5 (S. cerevisiae)
 guanine nucleotide binding protein (G protein), gamma 3
 putative c-Myc-responsive
 interleukin 8
 TJB protein
 PTX7 protein tyrosine kinase 7
 CGI-127 protein
 cathepsin L

neuroglobin
 PET12-like (yeast)
 Consensus includes gp:X81637.1 /DEF=H.sapiens clathrin light chain b gene, /FEA=mRNA /DB_XREF=gi:963046 /UG=Hs.73919 clathrin, light polypeptide (Lcb)
 cytochrome P450, subfamily 1E (ethanol-inducible)
 splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate splicing factor)
 KIAA0494 gene product
 protocadherin 9
 IQ motif containing GTPase activating protein 1
 sigma receptor (SR31747 binding protein 1)
 Consensus includes gp:AA906578 /FEA=EST /DB_XREF=gi:3042164 /DB_XREF=est:ot2206.x1 /CLONE=IMAGE:1524202 /UG=Hs.23158 ESTs
 solute carrier family 25 (mitochondrial carrier, peroxisomal membrane protein, 34kD), member 17
 Consensus includes gp:AI967961 /FEA=EST /DB_XREF=gi:5764779 /DB_XREF=est:wu11008.x1 /CLONE=IMAGE:2516655 /UG=Hs.26837 Homo sapiens mRNA; cDNA [AI967961
 phosphatidylinositol (4,5) bisphosphate 5-phosphatase, A
 ZYG homolog
 hypothetical protein FLJ20224
 glypican 1
 NAD(P)H dehydrogenase, quinone 1
 antigen identified by monoclonal antibody Ki-67
 BCL2/adenovirus E1B 19kD interacting protein 3-like
 RAN binding protein 9
 hypothetical protein FLJ23393
 nuclear transcription factor Y, alpha
 vacuolar protein sorting 41 (yeast)
 zinc finger protein, X-linked
 calcium channel, voltage-dependent, alpha 1G subunit
 hypothetical protein FLJ20039
 integrin, beta 2 (antigen CD18 (p95), lymphocyte function-associated antigen 1; macrophage antigen 1 (mac-1) beta subunit)
 homo box C4
 anaplastic lymphoma kinase (K1-1)
 hypothetical protein FLJ23416
 plasminogen-like
 jumonji homolog (mouse)
 oligodendrocyte myelin glycoprotein
 hypothetical protein FLJ11175
 delta-like homolog (Drosophila)
 netrin 1
 beta-transducin repeat containing
 ectodermal dysplasia 1, anhidrotic
 KIAA1111 protein
 hypothetical protein FLJ22060
 leucine-rich repeat protein, neuronal 1
 K01391 B subtilis TrpE protein, TrpD protein, TrpC protein corresponding to nucleotides 1883-4400 of K01391 [-5, -M, -3 represent transcript regions 5 prime, Middle, and 3 K01391
 hypothetical protein R29124_1
 DKFZP566H073 protein
 fibroblast growth factor receptor 1 (fms-related tyrosine kinase 2, Pfeiffer syndrome)
 gp:NM_005995.1 /DEF=Homo sapiens T-box 10 (TBX10) mRNA, /FEA=mRNA /GEN=TBX10 /PROD=T-box 10 /DB_XREF=gi:5174708 /UG=Hs.203847 T-box 10 /FL=gp:NM_005995
 Consensus includes gp:AU151793 /FEA=EST /DB_XREF=est:AU151793 /CLONE=NT2RP2008115 /UG=Hs.157078 Homo sapiens cDNA FLJ127
 hypothetical protein FLJ12409
 macrophage receptor with collagenous structure
 Consensus includes gp:AI207120 /FEA=EST /DB_XREF=gi:3765792 /DB_XREF=est:qf59a09.x1 /CLONE=IMAGE:1754344 /UG=Hs.246129 glutathione peroxidase 5 (epid AI207120
 heat shock 105kD
 Consensus includes gp:AW167793 /FEA=EST /DB_XREF=est:qf56607.x1 /CLONE=IMAGE:2632333 /UG=Hs.164038 Homo sapiens AKAP350C n AW167793
 hypothetical protein FLJ20321
 hypothetical protein MGCA309
 NM_021257
 NM_004584
 X81637
 AF182278
 M69040
 BC002525
 AI524125
 NM_003870
 AF228604
 AA906578
 AL048784
 AI967961
 AI935720
 NM_006336
 NM_017719
 NM_002081
 BC000906
 AU147044
 AF060922
 AA191426
 NM_024639
 AL031776
 NM_014396
 X59740
 AB012043
 NM_017635
 NM_000211
 AW298598
 U66559
 AK027069
 NS6120
 NM_004973
 NM_002544
 NM_018349
 U15979
 NM_004822
 AF101784
 AF061191
 AW249934
 NM_024612
 AI631881
 NM_024892
 NM_025105
 NM_008770
 NM_006770
 NM_006844
 NM_017786
 NM_024115

FIGURE 4C.

Gene	NCBI Accession Number	FOLD INCREASE
gp.L23516.1 /DEF=Human Ig rearranged gamma-chain, V-DP4-IH6, complete cds. /FEA=mRNA /DB_XREF=gi:385218 /FL=gb:L23516.1	L23516	43.37
ubiquitin carrier protein	NM_0114501	41.83
glyceraldhyde-3-phosphate dehydrogenase	M33197	41.07
Lysosomal-associated multispanning membrane protein-5	NM_006762	37.81
ribosomal protein L29	NM_009992	37.44
gp.L23518.1 /DEF=Human Ig rearranged gamma-chain, V-DP1-IH4b, complete cds. /FEA=mRNA /DB_XREF=gi:385220 /FL=gb:L23518.1	L23518	37.23
immunoglobulin heavy constant mu	U00139	33.58
adaptor-related protein complex 2, mu 1 subunit	NM_004068	32.97
heat shock 90D protein 1, beta	A218219	32.61
macrophage migration inhibitory factor (glycosylation-inhibiting factor)	NM_002415	30.90
lentin, light polypeptide	B538364	30.50
amino acid decarboxylase antizyme 1	A709094	30.15
Consensus includes gb:L48784 /DEF=OSO Homo sapiens cDNA, FEA=mRNA /DB_XREF=gi:1066715 /UG=Hs.182426 ribosomal protein 32	L48784	30.09
endoxin 1, (alpha)	U88968	28.76
ribosomal protein L29	BF683426	28.60
ribosomal protein L16a	NM_009979	28.20
chaperonin containing TCP1, subunit 7 (eta)	NM_009980	28.19
immunoglobulin heavy constant mu	NM_006429	27.91
MYT1 mitogen-activated protein kinase-2 (S. cerevisiae)	B3340548	27.05
protein 1	NM_001536	25.89
precollagen-protein, 2-oxoglutarate 4-dehydrogenase (proline 4-hydroxylase), beta polypeptide (protein disulfide isomerase; thyroid hormone binding protein p55)	NM_005022	24.23
precollagen-protein, 2-oxoglutarate 4-dehydrogenase (proline 4-hydroxylase), beta polypeptide (protein disulfide isomerase; thyroid hormone binding protein p55)	NM_000918	23.95
Consensus includes gb:D84143.1 /DEF=Human immunoglobulin (mAb59) light chain V region mRNA, partial sequence. /FEA=mRNA /PROD=immunoglobulin light chain V region /DB_XREF=IMMUNOGLOB	D84143	23.56
Consensus includes gb:A249377.1 /DEF=Homo sapiens partial mRNA for human Ig lambda light chain variable region, clone MB91 (331 bp). /FEA=mRNA /GEN=IGLV /PROD=immunoglob	A249377	22.30
US influenza-associated Sn-RNA protein LSm7	NM_016199	21.91
polymerase (DNA directed), delta 2, regulatory subunit (SNU2)	NM_006230	21.62
ribosomal protein L8	NM_000973	21.19
transketolase (Nernicke-Korshakoff syndrome)	BF686840	21.11
gp.L235092.1 /DEF=Homo sapiens mRNA for single-chain antibody, complete cds. /FEA=cDS /PROD=immunoglobulin /DB_XREF=gi:3090435 /UG=Hs.249245 Homo sapiens mRNA for sin	A0225092	20.48
T-cell leukemia/lymphoma 1A	B0001173	19.28
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	A02240	19.19
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	A0910614	19.08
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	B0005533	18.47
gp.L07950.1 /DEF=Homo sapiens H1C class 1 HLA B71 mRNA, complete cds. /FEA=cDS /GEN=HLA-B /PROD=HLA B71 /DB_XREF=gi:307236 /FL=gb:L07950.1	L07950	17.83
glyceraldhyde-3-phosphate dehydrogenase	M33197	17.71
ADMT7 mitochondriome maintenance deficient 7 (S. cerevisiae)	A2729900	17.26
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	NM_003751	16.97
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	NM_002654	16.81
pyruvate kinase, muscle	M87338	16.75
regulation factor 5 (activator 1) 2 (GND)	BF689355	16.61
glyceraldhyde-3-phosphate dehydrogenase	NM_003132	16.57
serine dehydratase	NM_005466	16.27
RNA polymerase II transcriptional regulation mediator (Med6, S. cerevisiae, homolog of)	NM_006423	16.24
ubiquitin-conjugating enzyme E2 (UBC9 homolog, yeast)	AF574664	16.05
ribosomal protein L13	NM_004140	15.99
Consensus includes gb:D84140.1 /DEF=Human immunoglobulin (mAb56) light chain V region mRNA, partial sequence. /FEA=mRNA /PROD=immunoglobulin light chain V region /DB_XREF=IMMUNOGLOB	D84140	15.89
CD22 antigen	552783	15.89
neutrophil cytosolic factor 4 (GND)	NM_013416	15.70
small EDRB-RN factor 2	NM_005770	15.52
glyceraldhyde-3-phosphate dehydrogenase	BE561479	15.36
transketolase (Nernicke-Korshakoff syndrome)	L12711	15.06
ATP synthase, H+ transporting, mitochondrial F1 complex, delta subunit	BF798517	14.96
CSG-135 protein	NM_016068	14.94
glucose phosphate isomerase	NM_000175	14.88
pain, desmosome associated protein	AF122222	14.86
actin, beta	X00351	14.30
Consensus includes gb:A254422 /DEF=Homo sapiens SHAR44 isoform (SHAR44) gene, complete cds, alternatively spliced /FEA=cDS_2 /DB_XREF=gi:10946127 /UG=Hs.78202 SWISN	A254422	14.20
lacton, derived from (11:2:16) malignant liposarcoma	NM_004960	14.10
ribosomal protein S5	NM_001009	13.79
polymerase (RNA) II (DNA directed) polypeptide 1 (13.3A)	B6335629	13.39

heat shock 90kD protein 1, beta	AF275319	13.32
nucleoside sensitive element binding protein 1	BC596374	13.25
ATP synthase, H ⁺ -transporting, mitochondrial F1 complex, delta subunit	NM_001687	13.24
edukaryotic translation initiation factor 3, subunit 8 (110kD)	AA679705	13.07
immunoglobulin heavy constant mu	BC001872	13.03
actin, beta	X00351	12.99
serine hydroxymethyltransferase 2 (mitochondrial)	NM_005412	12.93
polymerase (RNA) II (DNA directed) polypeptide F (25kD)	AA54759	12.83
maternal 610 transcript	NM_003910	12.80
major histocompatibility complex, class I, B	083043	12.59
major histocompatibility complex, class I, B	NM_015965	12.48
cell death-regulatory protein 6 (H19)	AL545523	12.22
proteasome (prosome, macropain) 26S subunit, ATPase, 3	AF100752	12.15
velocin-containing protein	NM_005804	11.98
nuclear RNA helicase, DEAD variant of DEAD box family	NM_003752	11.83
edukaryotic translation initiation factor 3, subunit 8 (110kD)	U34690	11.81
coronin, actin binding protein, 1A	NM_021107	11.76
mitochondrial ribosomal protein S12	NM_001280	11.67
cold inducible RNA binding protein	NM_016222	11.65
DEAD-box protein abstract	NM_020470	11.52
putative transmembrane protein; homolog of yeast Gaj1 membrane protein Yrip (Yip1p-interacting factor)	NM_002707	11.46
protein phosphatase 1c (formerly 2C), magnesium-dependent, gamma isoform	574639	11.35
immunoglobulin heavy constant mu	BC035989	11.32
acid phosphatase 1, soluble	NM_015380	11.29
CD451 protein	AL080089	11.11
ATP synthase, H ⁺ -transporting, mitochondrial F0 complex, subunit c (subunit 9), isoform 1	AW150953	11.06
7-dehydrocholesterol reductase	NM_001715	11.05
8 lymphoid tyrosine kinase	NM_005998	10.96
chaperonin containing TCP1, subunit 3 (gamma)	AF067524	10.85
cell division cycle 2 like 2	D25643	10.84
dehydrodiphenylphosphoglyoxalate-protein glycosyltransferase	BC000125	10.80
transforming growth factor, beta 1	NM_003363	10.71
dehydrocyclochrome c reductase core protein 1	NM_004542	10.61
MDH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3 (90D, B9)	NM_016034	10.58
mitochondrial ribosomal protein S2	AL545982	10.54
chaperonin containing TCP1, subunit 2 (beta)	AC004692	10.40
ribosomal protein L18a	AW402635	10.24
polymerase (RNA) II (DNA directed) polypeptide 1 (13.3kD)	NM_002201	10.17
interferon stimulated gene (20kD)	M24668	10.01
gpM24668.1 (DEF=Human Ig gamma chain V-region mRNA (C-D-JH4), complete cds. /FEA=nmRNA /GEN=IGH /DB_XREF=gi:185198 /FL=gb:M24668.1	NM_002812	9.92
proteasome (prosome, macropain) 26S subunit, non-ATPase, 8	AL078633	9.84
proteasome (prosome, macropain) subunit, alpha type, 7	BE239729	9.82
ribosomal protein S19	A0297586	9.71
major histocompatibility complex, class II, DR beta 5	AA284075	9.67
hesien 2 (60-70kD)	BE747342	9.65
similar to HYPOTHETICAL 34.0 XLA PROTEIN Z095.3 IN CHROMOSOME IV	NM_021154	9.50
phosphoserine aminotransferase	BE560461	9.50
sukla carrier family 2 (facilitated glucose/fructose transporter), member 5	D89646	9.48
mids homolog 6 (L. coli)	U02824	9.44
gpU02824.1 (DEF=Homo sapiens HLA class I heavy chain (HLA-Cw*1701) mRNA, complete cds. /FEA=CD5 /GEN=HLA-Cw*1701 /PROD=HLA class I heavy chain /DB_XREF=gi:1575443	L42024	9.43
stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing protein)	BE86580	9.39
nucleosome assembly protein 1 like 4	NM_005969	9.31
proteasome (prosome, macropain) 26S subunit, non-ATPase, 4	A033605	9.29
ribosomal protein S19	NM_001022	9.29
replication factor C (activator 1) 2 (40kD)	NM_002914	9.28
edukaryotic translation initiation factor 4 gamma, 1	AF104913	9.27
glutathione S-transferase pi	NM_000852	9.15
signal sequence receptor, delta (translocin-associated protein delta)	NM_006280	9.05
VP52B protein	NM_016208	9.01
DUF2756/H182 protein	AK000022	8.99
6-phosphogluconolactonase	NM_012088	8.94
ribosomal protein S2	NM_002952	8.88
protein phosphatase 1, regulatory subunit 7	BE718769	8.85

ribosomal protein S15	NM_001018	8.78
translocase of inner mitochondrial membrane 13 homolog 8 (yeast)	NM_012458	8.77
endothelial differentiation-related factor 1	AB02262	8.75
T cell receptor beta locus	M15564	8.61
hypothetical protein R33729_1	AC005339	8.53
ubiquitin C	M26880	8.53
major histocompatibility complex, class II, DR alpha	M60333	8.52
non-FOU-domain-containing, octamer-binding	L14599	8.52
major histocompatibility complex, class II, DR alpha	M60334	8.36
DNA replication factor	AF321125	8.34
PRP8 pre-mRNA processing factor 8 homolog (yeast)	NM_006445	8.28
SER1 homolog (S. pombe)	NM_006109	8.28
adrenyctic translation elongation factor 1 alpha 1	AL035687	8.27
KIA0217 protein	BC033881	8.24
major histocompatibility complex, class II, DR beta 1	U65585	8.20
ubiquitin-activating enzyme E1 (A15ST and BMT5 temperature sensitivity complementing)	NM_003334	8.14
polymerase (RNA) III (DNA directed) polypeptide K (12.3 kDa)	NM_016310	8.11
ATP synthase, H ⁺ -transporting, mitochondrial FO complex, subunit f, isoform 2	NM_004889	8.06
glutathionyl-HNA synthetase	NM_005051	8.04
HLA-B associated transcript 1	NM_004640	8.01
Consensus includes gpAF043584.1 /DEF=Homo sapiens clone ASHmg1-b1 immunoglobulin lambda chain VJ region, (J6L) mRNA, partial cds. /FEA=mRNA /GEN=HL /PROD=immunoglobul	AF043584	8.00
ribosomal protein S14	AF116710	7.98
interferon regulatory factor 4	NM_002460	7.94
T cell receptor beta locus	AF043179	7.91
HLA-B associated transcript 3	BC028444	7.90
bone marrow stromal cell antigen 2	NM_004335	7.84
integrin beta 4 binding protein	AF022229	7.76
putative breast adenocarcinoma marker (32kb)	NM_014453	7.65
polymerase (RNA) II (DNA directed) polypeptide L (7.6kD)	BC005903	7.63
hypothetical protein PR01847	AF119855	7.60
D123 gene product	NM_006023	7.54
actin binding LIM protein 1	NM_006720	7.53
adrenyctic translation initiation factor 3, subunit 2 (beta, 36kD)	U35764	7.47
T cell receptor beta locus	AL591122	7.45
tubulin, beta, 4	AL569749	7.39
customer protein complex, subunit epsilon	NM_007263	7.35
chromosome 14 open reading frame 3	NM_012111	7.35
major histocompatibility complex, class II, DP alpha 1	M27487	7.30
mitochondrial ribosomal protein S34	NM_023936	7.28
Hepatitis delta antigen-interacting protein A	NM_006848	7.27
pepdylprolyl isomerase F (cyclophilin F)	NM_005729	7.26
nuclear RNA export factor 1	BC004904	7.24
ubiquitin C	AB009010	7.21
transmembrane protein 4	BC001027	7.20
amyloid beta (A4) precursor-like protein 2	AW001847	7.19
hypothetical protein LU20113	AL523776	7.19
GCN1 general control of amino-acid synthesis 1-tila 1 (yeast)	AB91055	7.16
IMP2 non-histone chromosome protein 2-tila 1 (S. cerevisiae)	AF155235	7.15
adrenyctic translation initiation factor 4A, isoform 1	NM_001416	7.13
macrophage erythroblast attach	NM_005582	7.11
chemokine (C-X-C motif), receptor 4 (hsn)	AF348491	7.10
calreticulin	AD000092	7.09
ancient ubiquitous protein 1	NM_012103	7.07
enoyl-CoA hydratase	AF072405	7.06
putative human HLA class II associated protein 1	BE562032	7.05
suppression of tumorigenicity 14 (colon carcinoma, metastase, epithin)	NM_021978	7.05
proteasome (prosome, macropain) 26S subunit, non-ATPase, 3	NM_002809	7.02
GDP-mannose 4,6-dehydratase	NM_001500	7.01
transcription factor Dp-1	NM_007111	7.01
v-src myeloblastosis viral oncogene homolog (v-src) like 2	NM_002466	6.90
origin recognition complex, subunit 5-tila (yeast)	AF081459	6.89
Consensus includes gpAF005447.1 /DEF=Homo sapiens MHC class II antigen (DRB6) mRNA, HLA-DRB6-0201 allele, sequence. /FEA=mRNA /DB_XREF=g:5915893 /LG=Hs.167385 Homo	AF005487	6.89
splicing factor, arginine/serine-rich 5	AW084582	6.88

Consensus includes G08E905165 (FEA=EST /DB_XREF=gi:11717184 /DB_XREF=est:601186685T1 /CLONE=HMAE2959580 /UG=Hs. 100613 phospholipase C, beta 3, neighbor pseudogen	BE305165	6.88
ATP binding cassette, sub-family F (ABC2), member 2	NM_005692	6.87
adenoviral protein complex 2, sigma 1 subunit	NM_021575	6.86
U6 snRNA-associated Sm-like protein	NM_012321	6.86
unknown	NM_016209	6.85
major histocompatibility complex, class I, F	AW514210	6.84
ribosomal protein S2	A183766	6.82
HSV1 binding protein	NM_018694	6.81
CD75A antigen (immunoglobulin-associated alpha)	NM_001783	6.81
major histocompatibility complex, class II, DR beta 4	NM_021883	6.80
FTS66 binding protein precursor	NM_016594	6.77
eukaryotic translation initiation factor 3, subunit 4 (delta, 44KD)	BC007333	6.77
ADP-ribosylation factor 1	AI580004	6.76
hypothetical protein	NM_016459	6.74
anaphase promoting complex subunit 5	BC001081	6.74
U6 snRNA-associated Sm-like protein	AI12507	6.69
eukaryotic translation elongation factor 2	NM_001961	6.68
hypothetical protein M624675	AL118502	6.67
guanylate kinase 1	BC068249	6.66
SRP19 upstream reading frame	NM_022804	6.64
mitochondrial ribosomal protein L23	A832239	6.62
RNA, U2, small nuclear	BC003629	6.58
membrane-spanning 4-domains, subfamily A, member 1	X12530	6.57
head shock protein 75	NM_016292	6.51
transglutinin 2	NM_003564	6.50
trypophanyl-tRNA synthetase	M61715	6.47
actin, gamma 1	AL567820	6.46
eukaryotic translation initiation factor 4A, isoform 1	BC066210	6.45
guanine nucleotide binding protein (G protein), beta polypeptide 1	A741124	6.45
ribosomal protein, large, P0	A953822	6.43
ribonucleotide reductase M1 polypeptide	A692974	6.42
proteasome (prosome, macropain) 26S subunit, non-ATPase, 2	NM_002808	6.42
CD27-binding (Shc) protein	NM_006427	6.38
small nuclear ribonucleoprotein polypeptides B and B1	J04564	6.36
membrane-spanning 4-domains, subfamily A, member 1	BC002807	6.33
ribosomal protein, large, P0	NM_001002	6.32
ribosomal protein, large, P0	BC003863	6.30
sub4D	AI109955	6.28
ribosomal protein, large, P0	BC003655	6.28
mitochondrial ribosomal protein S16	NM_016065	6.26
CD27-binding (Shc) protein	A5033111	6.23
steroid regulatory element binding transcription factor 2	NM_004599	6.23
CDWS2 antigen (CDPAT11 antigen)	NM_001803	6.21
ribosomal protein L10	NM_006013	6.21
cytochrome c oxidase subunit IV isoform 1	NM_001861	6.19
ribosomal protein S3	U14990	6.17
hydroxymethyl-Coenzyme A dehydrogenase/3-ketoacyl-Coenzyme A thiolase/enoyl-Coenzyme A hydratase (trifunctional protein), alpha subunit	A972144	6.15
neurospiral cytosolic factor 4 (NCF4)	NM_006631	6.15
NADH dehydrogenase (ubiquinone) flavoprotein 1 (51kD)	A792131	6.14
ascorbate 2, mitochondrial	NM_001098	6.11
ribosomal protein S26	NM_001029	6.10
HLA-B histocompatibility antigen, class I, E	A7226990	6.10
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 4	A744900	6.08
DNA segment on chromosome 19 (unique) 1177 expressed sequence	A4389278	6.05
proteasome (prosome, macropain) subunit, beta type, 7	NM_002799	6.05
lymphocyte-specific protein tyrosine kinase	NM_005356	6.04
seiy-dRNA synthetase	NM_006513	6.04
v-pes-1 Yamaguchi sarcoma viral related oncogene homolog	M79321	6.03
HSPK142 protein	BC006244	6.02
replication protein A1 (70kD)	NM_002945	5.99
ubiquitin-conjugating enzyme E26 2 (UBK7 homolog, yeast)	B6395660	5.98
membrane component, chromosome 11, surface marker 1	B6258784	5.96

H1 histone family, member X	NM_006026	5.95
lysosomal-associated multispinning membrane protein-5	AS99086	5.95
PTD008 protein	NM_016145	5.95
poly(IG) binding protein 1	U24223	5.94
polyglutamine binding protein 1	A0041836	5.93
active homology 2 (Trospolia)	BC000422	5.93
MAOS box transcription enhancer factor 2, polypeptide B (myocyte enhancer factor 2B)	NM_005919	5.92
translocase of inner mitochondrial membrane 44 homolog (yeast)	NM_006351	5.92
intercellular adhesion molecule 2	NM_000873	5.91
bromodomain-containing 2	D42040	5.91
lysosomal-associated membrane protein 1	NM_005561	5.91
MDH dehydrogenase (ubiquinone) 1 beta subcomplex, 8 (19kD, ASH1)	NM_005004	5.87
actin related protein 2/3 complex, subunit 2 (34 kD)	A2739893	5.84
Consensus includes gb:83344495 /FE=EST /DB_XREF=gi:13140933 /DB_XREF=est:602461128f1 /CLONE=IMAGE-4577718 /UG=Hs.228201 ESTs, Moderately similar to AF118094.25 PF	NM_004135	5.80
bicucitrate dehydrogenase 3 (MDH+3) gamma	BC334495	5.80
mitochondrial ribosomal protein L9	NM_007108	5.80
transcription elongation factor 8 (SIII), polypeptide 2 (18kD, elongin B)	A670847	5.80
hypothetical protein	NM_014145	5.77
HSP274 protein	M60261	5.77
APF1 nuclease (multifunctional DNA repair enzyme)	NM_016185	5.75
hematological and neurological expressed 1	NM_025205	5.75
hypothetical protein DKE24248185	AF047695	5.75
calcaryotic translation initiation factor 4E-like 3	NM_001640	5.74
N-acylfluoroyl-peptide hydrolase	AA52514	5.73
QMA0746 protein	A0001527	5.72
calpain, small subunit 1	AF147209	5.69
interleukin enhancer binding factor 3, 90kD	NM_006773	5.67
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 18 (Myc-regulated)	AL049650	5.67
hypothetical protein M6C10715	AF034607	5.66
chloride intracellular channel 1	NM_000626	5.64
CD79b antigen (immunoglobulin-associated beta)	NM_004493	5.64
hydroxyacyl-Coenzyme A dehydrogenase, type II	BE871379	5.62
zinc finger protein 207	AC065011	5.61
ribosomal protein S28	A0285970	5.61
synactin 16	NM_012336	5.61
nuclear prelamin A recognition factor	NM_000101	5.60
cytochrome b-245, alpha polypeptide	NM_006409	5.60
actin related protein 2/3 complex, subunit 1A (41 kD)	NM_021126	5.57
metacasparynsulfate sulfatransferase	BF979419	5.57
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lymphocyte cytosolic protein 1 (L-plastin)	NM_004551	5.56
MDH dehydrogenase (ubiquinone) Fe-S protein 3 (30kD) (MDH-coenzyme Q reductase)	NM_004596	5.55
small nuclear ribonucleoprotein polypeptide A	BF342707	5.55
pre-mRNA processing factor 31 homolog (yeast)	NM_002708	5.54
protein phosphatase 1, catalytic subunit, alpha isoform	NM_003321	5.54
Ta translation elongation factor, mitochondrial	AA514622	5.52
QMA0518 gene product	U34994	5.51
protein kinase, DNA-activated, catalytic polypeptide	M12679	5.50
Ccr1 antigen	NM_014225	5.49
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), alpha isoform	NM_003406	5.49
tyrosine 3-monooxygenase/tyrosylphosphatase 5-monooxygenase activator protein, zeta polypeptide	NM_005273	5.45
guanine nucleotide binding protein (6 protein), beta polypeptide 2	AF029750	5.43
TAP binding protein (tapasin)	NM_002394	5.42
soluble carrier family 3 (members of dibasic and neutral amino acid transport), member 2	NM_003145	5.41
signal sequence receptor, beta (translocase-associated protein beta)	U17496	5.36
proteasome (prosome, macropain) subunit, beta type, 8 (large multifunctional protease 7)	NM_016337	5.33
BDG5	NM_000581	5.29
glutathione peroxidase 1	BF570122	5.28
QMA0123 protein	NM_004111	5.27
flag structure-specific endonuclease 1	BC005876	5.27
ATPase, H+-transporting, lysosomal (vacuolar proton pump) 21kD	NM_017528	5.27
putative methyltransferase	U23946	5.26
RNA binding motif protein 5		

GTP cyclohydrolase 1 feedback regulatory protein	NM_005258	5.25
12A histone family, member O	A313324	5.25
hypothetical protein B32184_1	BC001648	5.25
larnesyl-diphosphate larnesyltransferase 1	BC003573	5.25
phycocyanin, synaptic 2	NM_004668	5.22
peptide/protein isomerase E (cyclophilin E)	AF042386	5.21
PKS6 binding protein 1A (123D)	BC005147	5.19
guanine nucleotide binding protein (G protein), beta polypeptide 2-like 1	NM_006098	5.15
lymphocyte-specific protein tyrosine kinase	U07236	5.13
MO15 melanosome maintenance deficient 5, cell division cycle 46 (S. cerevisiae)	NM_006739	5.12
homo box A1	AC004079	5.11
HLA-B associated transcript 3	NM_004639	5.11
tubulin alpha 6	BC005946	5.10
endothelinase 6	NM_004435	5.10
similar to RDE31 cDNA Z310040617 gene	BF972185	5.09
protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific, extinguisher 1)	M18468	5.09
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protein disulfide isomerase related protein (calcium-binding protein, intestinal-related)	BC000425	5.07
proteasome (prosome, macropain) subunit, beta type, 3	NM_002795	5.07
T cell receptor alpha locus	M12423	5.06
MCOT mitochondrion maintenance deficient 7 (S. cerevisiae)	D55716	5.06
nuclear distribution gene C (A. nidulans) homolog	AF241788	5.05
T cell receptor alpha locus	L34703	5.04
HLA-B associated transcript 3	BC003133	5.02
ATPase, H+-transporting, lysosomal (vacuolar proton pump), member D	AL566172	5.00
actin, gamma 1	AF145192	4.99
histon-H2a 4	AC002301	4.98
haryphatin (limpstatin) beta 3	NM_002271	4.98
translocase of inner mitochondrial membrane 10 homolog (yeast)	NM_012456	4.98
bacterial 10P repeat-containing 5 (survivin)	A8028869	4.98
stress-induced-phosphoprotein 1 (Hsp70/Hsp90-organizing protein)	AL553320	4.97
ribosomal protein, large, P1	NM_001003	4.97
myosin 1D	A4621962	4.97
2q28.2000bp sequence contig, ORF	BE576218	4.96
GDP-asease phosphatophylase A	NM_013335	4.96
casein kinase 2, beta polypeptide	NM_001320	4.91
proteasome (prosome, macropain) 35S subunit, non-ATPase, 4	U51007	4.90
culin 1	NM_003592	4.90
Hurler's interacting protein 8	AF049103	4.90
HLA-E histocompatibility antigen, class I, C	M90684	4.87
solute carrier family 7 (cationic amino acid transporter, y+ system), member 5	A8018009	4.86
meic fringe homolog (Drosophila)	A738965	4.85
RNA-binding protein (leucostigminic)	NM_016732	4.85
heat shock 70MD protein 4	BC002526	4.84
SRH/SHF related, matrix associated, actin dependent regulator of chromatin, subfamily f, member 1	AF231056	4.84
leucine rich repeat (in FLU) interacting protein 1	NM_004735	4.83
Rho GDP dissociation inhibitor (GDI) alpha	NM_004309	4.83
adaptor-related protein complex 2, sigma 1 subunit	BC006337	4.82
lymphoblastin beta (TNF superfamily, member 3)	NM_002341	4.81
cytochrome c oxidase subunit Vb	NM_001862	4.81
chromosome 11 open reading frame2	NM_013265	4.79
NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 7 (14.5D, 8.14.5a)	NM_005001	4.78
RNA4905 protein	AA502912	4.77
splicing factor, arginine/serine-rich 1 (splicing factor 2, alternate splicing factor)	NM_006924	4.77
CD37 antigen	NM_001774	4.77
tubulin alpha 6	BC004949	4.74
block of proliferation 1	B6491842	4.74
peroxiredoxin 1	L19164	4.73
thymopressin	AF113682	4.72
cyclic-dependent kinase 4	NM_000075	4.71
ribosomal protein L13a	BC001675	4.70
T cell receptor alpha locus	M15565	4.69
PAI-1 mRNA-binding protein	BC003049	4.66

DNF2P4341813 protein	B6168666	4.14
c-2 and 5-phase expressed 1	BF305380	4.14
ribosomal protein L27a	NM_000990	4.14
neutrophil cytosolic factor 1 (47kD, chronic granulomatous disease, autosomal 1)	NM_000265	4.13
serum/glucocorticoid regulated kinase-like	NM_013257	4.13
FGS6 binding protein 1A (120k)	BC001002	4.13
phosphoglucosyl dehydrogenase	NM_002631	4.13
non-POU-domain-containing, octamer-binding	BC003129	4.12
RNA binding motif protein 10	AL137421	4.12
serpinyl antipeptidase	NM_005724	4.12
tetraspin 3	NM_005724	4.12
Erk1g sarcoma breakpoint region 1	BC006327	4.10
ret finger protein	AF230394	4.10
ribonuclease S precursor	NM_003730	4.09
tumor protein D52	BC389015	4.09
Consensus includes gb:46537190 /FEA=EST /DB_XREF=gi:13528922 /DB_XREF=IMAGE:4690079 /UG=Hs.111334 ferritin, light polypeptide	BC537190	4.08
BCL2-interacting killer (apoptosis-inducing)	NM_001197	4.08
RUM310 gene product	BC001404	4.07
accessory proteins BMP31/AMF29	NM_005745	4.06
proteasome (prosome, macropain) 26S subunit, non-ATPase, 9	NM_002813	4.05
neutrophil cytosolic factor 1 (47kD, chronic granulomatous disease, autosomal 1)	AW072388	4.05
butyrophilin, subfamily 3, member A2	BC002832	4.04
B-cell associated protein	NM_007273	4.03
ATPase, H ⁺ transporting, lysosomal (vacuolar proton pump), beta polypeptide, 56/58kD, isoform 2	NM_001693	4.02
mitochondrial ribosomal protein S7	NM_015971	4.01
major histocompatibility complex, class I, E	NM_005516	4.00
mitogen-activated protein kinase kinase 3	AA703381	4.00
RUM4599 protein	BC002327	3.99
protein phosphatase 1, regulatory subunit 7	NM_002712	3.99
RUM0729 protein	AW502434	3.98
thyroid autoantigen 70kD (Mu antigen)	NM_001469	3.97
proteasome (prosome, macropain) subunit, beta type, 9 (large multifunctional protease 2)	NM_002800	3.97
hypothetical protein MGC5585	NM_024057	3.96
polymyoma (RNA) II (DNA directed) polypeptide 1 (14.5kD)	AL037557	3.94
B-cell translocation gene 1, anti-proliferative	AL535380	3.94
hypothetical protein FLJ20859	NM_022734	3.93
ATP citrate lyase	U18197	3.92
GMS complex locus	AF064092	3.92
gb:M24665.1 /DEF=Human Ig rearranged H-chain V-region mRNA (c-DJH6), complete cds. /FEA=miRNA /GEN=IGHG /DB_XREF=gi:185200 /FL=gb:M24665.1	M24669	3.92
major histocompatibility complex, class I, E	M31183	3.91
insulin induced gene 1	NM_005542	3.91
Consensus includes gb:4001784 /FEA=EST /DB_XREF=gi:3202255 /DB_XREF=est:041 g06.s1 /CLONE=IMAGE:1619386 /UG=Hs.308332 ESTs. Highly similar to M42735 ribosomal protein	NM01784	3.91
Consensus includes gb:282202 /DEF=Human DNA sequence from clone RP1-34P24 on chromosome 22. Contains a pseudogene similar to ribosomal protein L35, E51s, S17s and G5s	Z82202	3.89
heterogeneous nuclear ribonucleoprotein R	NM_005826	3.89
platelet activating factor acetylhydrolase, isoform 1b, gamma subunit (29kD)	NM_002573	3.89
hypothetical protein	BC257762	3.88
cytochrome c oxidase subunit Vb	BC096229	3.88
DNF2P547E1010 protein	AF261137	3.88
interferon regulatory factor 3	NM_001571	3.87
postmitotic segregation increased 2-like 9	U38979	3.86
putative cyclin G1 interacting protein	NM_006349	3.86
GMS complex locus	NM_000516	3.86
cyclin-dependent kinase 5	NM_004935	3.86
hepatocellular carcinoma 1	NM_002149	3.86
24-dehydrocholesterol reductase	NM_014762	3.86
2',5'-bisphosphoryl synthetase 1 (40-46 kD)	NM_002534	3.85
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 30	NM_014966	3.85
CREBBP/EP300 inhibitory protein 1	AF274951	3.85
adenylate kinase 2	U39945	3.84
structure specific recognition protein 1	BE795648	3.84
Rho GTP dissociation inhibitor (GDI) alpha	BC005851	3.84
Consensus includes gb:304602 /DEF=Homo sapiens UBBP2 pseudogene for ubiquitin UB8 /FEA=CD5 /DB_XREF=gi:37582 /UG=Hs.247890 Homo sapiens UBBP2 pseudogene for ubiquitin	X04802	3.84
ubiquitin-conjugating enzyme E2H (UBK12 homolog, yeast)	NM_003969	3.83

adenosine deaminase	NM_000022	3.83
leucine 2 (60-70kD)	A0284075	3.82
autolytic translation initiation factor 2, subunit 3 (gamma, 52kD)	NM_001615	3.82
hypothetical protein P101847	AF119855	3.82
enhancer of invasion 10	NM_021178	3.82
HOX2 mitochromosome maintenance deficient 2, mitotin (S. cerevisiae)	NM_004526	3.81
WD repeat domain 1	U63131	3.81
OC37 cell division cycle 37 homolog (S. cerevisiae)	A8010427	3.81
WD repeat domain 1	BEF79367	3.81
A kinase (PKA) anchor protein 2	NM_006156	3.80
neural precursor cell expressed, developmentally down-regulated 8	NM_018835	3.80
dictyary receptor, family 1, subfamily K, member 1	AF088184	3.79
GMS complex locus	NM_006327	3.79
translocase of inner mitochondrial membrane 23 homolog (yeast)	NM_002412	3.79
O-6-methylguanine-DNA methyltransferase	NM_001402	3.79
autolytic translation elongation factor 1 alpha 1	NM_005324	3.79
H3 histone, family 3B (H3.3B)	BE551340	3.78
RNA974 protein	M21154	3.77
S-adenosylmethionine deacylase 1	NM_031213	3.76
hypothetical protein M625244	NM_012225	3.75
nucleoside binding protein 2 (RibD homolog, E. coli)	NM_001913	3.75
cat-dna 1, COAT1 displacement protein (Drosophila)	NM_006802	3.75
splicing factor 3a, subunit 3, 60kD	NM_022551	3.74
ribosomal protein S18	NM_003904	3.74
zinc finger protein 259	NM_013440	3.73
paired immunoglobulin-like receptor beta	NM_003600	3.73
serine/threonine kinase 15	AF098158	3.73
chromosome 20 open reading frame 1	NM_031286	3.72
SH3 domain binding glutamic acid-rich protein like 3	L49506	3.72
cyclin G2	NM_005722	3.71
ANP2 actin-related protein 2 homolog (yeast)	NM_003470	3.69
ubiquitin specific protease 7 (herpes virus-associated)	NM_001930	3.69
deoxyhypusine synthase	BC001149	3.69
serologically defined colon cancer antigen 16	AL031276	3.68
ribosomal protein L10	A336206	3.68
PRKX, apoptosis, WT1, regulator	U04627	3.67
hydroxyaryl-Coenzyme A thiolase/aryl-Coenzyme A thiolase (enoyl-Coenzyme A hydratase (bifunctional protein), alpha subunit	225433	3.66
gla-225433.1 / RFE-H sapiens protein-serine/threonine kinase gene, complete CDS. / FEA-mRNA / P100D=protein-serine/threonine kinase / DB_XREF=gi:405744 / FL=gb:225433.1	AI167775	3.66
HLA-G-related gene	NM_016645	3.65
mesenchymal stem cell protein DSC2	AW007161	3.65
SXT (sets determining region 1)-box 2	NM_007192	3.65
chromatin-specific transcription elongation factor, 140 kDa subunit	U62819	3.64
uncoupling protein 2 (mitochondrial proton carrier)	A1026905	3.63
leucine-3aa 5 (mitotic centrosome-associated kinesin)	NM_006152	3.63
lymphoid-restricted membrane protein	BC03574	3.63
T-cell leukemia/lymphoma 1A	NM_0071695	3.63
ATPase, H+-transporting, lysosomal (vacuolar proton pump) 42kD	NM_005483	3.61
chromatin assembly factor 1, subunit A (p150)	NM_005789	3.60
proteasome (prosome, macropain) activator subunit 3 (PA28 gamma; k)	BC000779	3.60
RNA1007 protein	NM_004877	3.59
gla maturation factor, gamma	BC001430	3.59
POP7 (processing of precursor, S. cerevisiae) homolog	A044773	3.59
breakpoint cluster region protein, uterine leiomyoma, 1; barrier to autointegration factor	NM_024292	3.59
ubiquitin-like 5	NM_030799	3.58
hypothetical protein NF140225	BC000924	3.58
IBR2	U26266	3.58
deoxyhypusine synthase	NM_005809	3.58
peroxiredoxin 2	BE731738	3.58
Consensus includes gb:BE731738 / DB_XREF=gi:10145730 / DB_XREF=est:60156815461 / KLINE=IMAGE:3842844 / U6=Hs_182937 peptidylprolyl isomerase A (cyclophilin A)	NM_003971	3.57
spasm associated antigen 9	A059865	3.57
HOX4 mitochromosome maintenance deficient 4 (S. cerevisiae)	AI136306	3.57
leucine receptor 1 (67kD, ribosomal protein 5A)	A4670344	3.57
villin 2 (cecr1)	NM_015871	3.57
zinc finger protein		

ubiquitin 8	204801	3.56
metastasis-associated 1-like 1	BE222618	3.56
hypothetical protein FLJ10597	NM_018150	3.54
similar to mouse G13 or D. melanogaster transcription factor 118	NM_012242	3.54
proboscidea (proscione, macroquin) activator subunit 3 (P428 gamma; K)	BC026894	3.53
intercellular adhesion molecule 3	NM_002162	3.53
hypothetical protein MGC10433	NM_024321	3.53
mitochondrial ribosomal protein S17	NM_015989	3.52
endothelial differentiation-related factor 1	AB022282	3.52
gamma-tubulin complex protein 2	NM_006659	3.51
sialyltransferase 1 [beta-galactoside alpha-2,6-sialyltransferase]	AV695711	3.51
transcription factor CP2	U03494	3.51
KIA0152 gene product	NM_014730	3.51
Fancet anemia, complementation group A	AF033279	3.51
high-mobility group 208	AF288679	3.50
zinc finger protein 313	AL031685	3.50
cellular protein elongation factor 1 gamma	AF119850	3.49
nuclear protein family A, member 3 (N/A) small nuclear RNPs	NM_018648	3.48
NADH dehydrogenase (ubiquinone) Fe-S protein 8 (Z3U0) (NADH-coenzyme Q reductase)	NM_002496	3.48
proteasome protein 2 (colonie epithelium-enriched)	NM_002668	3.48
p47	NM_016143	3.47
KIA0074 protein	D38553	3.47
nuclear distribution gene C (A. nidulans) homolog	NM_006600	3.46
cytochrome c-1	NM_001916	3.46
Consensus includes gb:574774.1 /REF=g599n(T) =OMT3-induced calcium influx regulator human, Jurkat J6 T cell line, mRNA Partial, 1605 nt. /FEA=CD5 /PROD=tyrosine kinase p59n(T)	374774	3.46
hypothetical protein FLJ10479	NM_018106	3.46
hypothetical protein FLJ12287 similar to semaphorins	NM_023367	3.45
apoptosis antagonizing transcription factor	AF083208	3.45
cellular protein elongation factor 1 alpha 1	NM_001403	3.44
ribosomal protein L15	AF279903	3.44
chromosome condensation 1	NM_001269	3.43
hypothetical protein FLJ10439	BC006351	3.43
gb:AT014272.1 /REF=Homo sapiens F55330 mRNA, complete cds. /FEA=mRNA GEN=F55330 /PROD=Homo sapiens F55330 /AV014272	NM_004158	3.43
Consensus includes gb:002723.1 /REF=Homo sapiens cDNA: FLJ23572 fs, clone UG12403. /FEA=mRNA /DB_JREF=gi:10440306 /UG=Hs.201593 Homo sapiens cDNA: FLJ23572 fs, d	U02725	3.43
kinase-like 8 (mitotic centrosome-associated kinesin)	U03743	3.42
SHC (Src homology 2 domain-containing) transforming protein 1	A091079	3.42
succinate dehydrogenase complex, subunit A, flavoprotein (Fp)	NM_004158	3.42
casein kinase 2, beta polypeptide	M30448	3.42
adenylate kinase 2	W02312	3.42
lysine 3-monooxygenase/hypophar 5-monooxygenase activation protein, theta polypeptide	NM_006826	3.42
glutamic pyruvate transaminase (alanine aminotransferase)	NM_005309	3.42
tricarboxylate isomerase 1	BF116254	3.42
breast carcinoma amplified sequence 4	NM_017843	3.42
ribosomal protein L10	AF057781	3.42
prolactin	NM_002634	3.42
zinc finger protein 183 (RING finger, CHK4 type)	BC008032	3.41
springstone-1-phosphatase	NM_030791	3.41
hypothetical protein MGC11256	NM_024324	3.41
nucleophosmin/nucleophosmin 3	NM_006993	3.40
hypothetical protein FLJ11856	AK021918	3.40
LIN and SH3 protein 1	NM_006148	3.40
ribonucleotide reductase M2 polypeptide	BC001886	3.40
casein kinase 1, delta	AV700224	3.39
forkhead box M1	NM_021953	3.39
endoxin 1, (alpha)	NM_001428	3.38
DNA binding protein 7	AL173325	3.38
general transcription factor IIC, polypeptide 3 (102kd)	NM_012086	3.38
postmitotic segregation increased 2-like 11	U38960	3.38
pyridoxal (pyridoxine, vitamin B6) kinase	NM_003681	3.37
hypothetical protein FLJ20340	NM_017773	3.37
alanyl-RNA synthetase	NM_001605	3.37
luciferase 1 (E. coli)	NM_003707	3.36
Consensus includes gb:86537255 /FEA=EST /DB_JREF=gi:13529117 /DB_JREF=est:602565318F1 /CLONE=IMAGE:4689748 /UG=Hs.82646 Dnal (Hsp40) homolog, subfamily B, member	86537255	3.36

Ras-4Tase activating protein SH3 domain-binding protein 2	NM_012297	3.36
SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily b, member 1	AK021419	3.35
hypothetical protein IMAGE3455200	NM_024006	3.34
coupling protein (actin filament) muscle Z-line, beta	NM_004930	3.34
Dual (Dpp40) homolog, subfamily C, member 8	AA843895	3.34
transportin-SR	NM_012470	3.34
cellular (p130cas) assembly 1-like	BC001663	3.33
Sigma3 synthase nuclear activator 1	BC004118	3.33
protein synthetase co-transcribed (bacterial homolog)	NM_007198	3.33
interleukin stimulator gene (IL20)	U88564	3.32
protease (prosome, macropain) subunit, beta type, 4	NM_002796	3.32
serine hydroxymethyltransferase 2 (mitochondrial)	AF150316	3.32
HPF2 non-histone chromosome protein 2-isoa 1 (S. cerevisiae)	NM_005008	3.32
serine/threonine kinase 12	AB011446	3.32
COP9 homolog	NM_006710	3.32
HSX150 protein	NM_014182	3.31
nuclear matrix protein HNP200 related to splicing factor PRP19	NM_014502	3.31
myosin, light polypeptide 6, alpha, smooth muscle and non-muscle	BE734356	3.31
ATX1 antioxidant protein 1 homolog (yeast)	NM_004045	3.31
MTX1 protein	NM_021640	3.31
hyphal-restricted membrane protein	U10485	3.30
PD RNA-binding protein	U03411	3.29
histidyl-RNA synthetase	NM_002109	3.29
hypothetical protein MG2650	NM_024108	3.29
adenine phosphoribosyltransferase	AA027724	3.28
BDJ23 homolog A (S. cerevisiae)	BF572938	3.28
g0AF274954.1 (DEF=Homo sapiens PNAS-29 mRNA, complete cds. /FEA=mRNA /PID=PNAS-29 /DB_XREF=gi:12751070 /UG=Hs.326791 Homo sapiens PNAS-29 mRNA, complete cds. /)	AF274954	3.28
hypothetical protein FUJ2549	NM_024644	3.28
immunoglobulin lambda locus	AF047245	3.27
hypothetical protein FUJ1305	NM_018386	3.27
HSPC002 protein	NM_015362	3.27
translin	AF35302	3.25
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	M65254	3.24
death associated protein 3	U18321	3.24
SHG-interacting protein	AF889564	3.24
hypothetical protein FLJ20552	NM_017876	3.23
uncharacterized hypothalamus protein HT009	NM_018470	3.23
mitochondrial ribosomal protein L12	NM_002949	3.23
memusidase, alpha, class 1A, member 1	NM_005907	3.23
lactone H2A17Z variant	BF343852	3.23
dihydrodipicolinate 5-acyltransferase (E2 component of pyruvate dehydrogenase complex)	J03866	3.23
ATP-dependent interferon responsive	NM_022371	3.23
nu25 homolog 5 (E. coli)	AF034759	3.23
X-prolyl aminopeptidase (aminopeptidase P)-like	NM_006523	3.22
villin 2 (catrin)	J05021	3.22
map kinase phosphatase-like protein MK-STYK	NM_016086	3.22
ricaphorin II	BC003560	3.22
RAM1608 protein	NM_024820	3.21
G3 protein	NM_016947	3.21
eukaryotic translation initiation factor 2B, subunit 2 (beta, 39kD)	NM_014239	3.21
nuclear transcription factor Y, gamma	U62296	3.20
Integral type I protein	BC000027	3.20
ATP synthase, H ⁺ -transporting, mitochondrial F1 complex, beta polypeptide	NM_001686	3.20
Consensus includes g0:AK025759.1 /DEF=Homo sapiens cDNA: FLJ22106 fs, clone HEP17675, /FEA=mRNA /DB_XREF=gi:10438374 /UG=Hs.323470 Homo sapiens cDNA: FLJ22106 fs, d	AK025759	3.20
dyx19c1, cytoplasmic, heavy polypeptide 1	AB002323	3.20
change and polyadenylation specific factor 1, 16kD subunit	NM_013291	3.20
nuclei (nucleoside diphosphate linked moiety X)-type motif 1	NM_002452	3.19
hypothetical protein FUJ0374	NM_018074	3.19
glutamic-carbaloic acid transaminase 2, mitochondrial (aspartate aminotransferase 2)	NM_002080	3.18
candidate tumor suppressor protein	NM_020381	3.17
RAM1049 protein	AF322111	3.17
Consensus includes g0:AL18510 /DEF=Human DNA sequence from clone RPS-858M22 on chromosome 20. Contains a pseudogene similar to 40S ribosomal protein S10, STSs and G55a /FEA=AL18510	AF322111	3.17
tumor suppressor deleted in oral cancer-related 1	NM_005851	3.16

phosphoserine isomerase 1	NM_000365	3.16
phosphoribosyl pyrophosphate synthetase 1	NM_002764	3.16
aldo-keto reductase family 1, member A1 (aldehyde reductase)	NM_006066	3.15
major histocompatibility complex, class II, DR beta 5	NM_002125	3.15
aryl hydrocarbon receptor interacting protein	AL558532	3.15
prostate tumor over expressed gene 1	AL046054	3.15
anaphase promoting complex subunit 5	BF967271	3.15
peptide/protein isomerase M (cyclophilin H)	NM_006347	3.14
major histocompatibility complex, class I, C	AB248336	3.14
adenylosuccinate lyase	NM_000026	3.13
adenylyl transferase initiation factor 3, subunit 9 [eta, 1184D]	U78525	3.13
DC12 protein	NM_020187	3.13
G4E binding protein	NM_014593	3.13
divalent cation tolerant protein (UTA)	AF210924	3.12
hypothetical protein	AA243508	3.12
NAOH dehydrogenase (ubiquinone) Fe-S protein 6 (13L0) (NAOH-coenzyme Q reductase)	NM_004553	3.12
Consensus includes gb:BE547674 /FEA=EST /DB_XREF=gi:9776319 /DB_XREF=est:601076255F1 /CLONE=IMAGE:3462325 /UG=Hs.204169 ESTs	BE547674	3.11
hypothetical protein similar to mouse aldehyde reductase 6 (mral)	AL096767	3.11
GMP12 for guanine triphosphate reductase isolog	NM_016576	3.11
vesicle-associated membrane protein 1 (Glypobrevin 1)	NM_016830	3.11
acetonamide nucleotide transferase	NM_012343	3.10
cell division cycle 2, G1 to S and G2 to M	NM_007186	3.10
hypothetical protein FLJ12660	NM_025152	3.10
BAB7, member RAS oncogene family	AK000826	3.10
nucleoside phosphorylase	NM_000270	3.10
sorting nexin 5	NM_014426	3.09
core histone macroH2A.2	NM_018649	3.09
adenylyl transferase initiation factor 4E binding protein 1	AB044548	3.08
ELAV (elavryne) lethal, abnormal vision, (Drosophila)-like 1 (Hu antigen B)	NM_001419	3.08
mitochondrial ribosomal protein S18B	BC053773	3.08
peccabdo homolog 1, containing BKT domain (zobretsh)	NM_014303	3.07
GTP dissociation inhibitor 2	NM_001494	3.07
lysine 3-monoxygenase/hypophan 5-monoxygenase activation protein, epsilon polypeptide	U28936	3.07
BD repeat domain 6	NM_018031	3.07
splicing factor, arginine/serine-rich 2, interacting protein	NM_004719	3.07
hypothetical protein FLJ20436	AK022732	3.07
glycoprotein	NM_004130	3.07
phosphoribosyl aminotransferase	AB89380	3.07
dyskeratosis congenita 1, dyskerin	NM_001363	3.07
cytoskeleton-associated protein 1	NM_001281	3.07
neural precursor cell expressed, developmentally down-regulated 5	AI91427	3.07
bleomycin hydrolase	NM_000386	3.06
DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 1	NM_004939	3.06
serpin-related receptor, L (DLR class) A repeats-containing	NM_003105	3.06
damage-specific DNA binding protein 1 (127AD)	L40326	3.06
ribosomal protein L15	Z97353	3.06
PP3111 protein	NM_022156	3.05
sorting nexin 1	AF05484	3.05
zinc finger protein 143 (clone pHZ-1)	AF05484	3.05
C-terminal binding protein 1	AF05484	3.05
hypothetical protein FLJ10989	AF05484	3.05
UBL domain-containing 2	NM_001328	3.05
ribosomal protein S2	NM_018292	3.05
B-cell CLL/lymphoma 6 (zinc finger protein 51)	AB30314	3.04
arx1 (bacterial) arx1-like transporter, ATP-binding, homolog 1	AF05484	3.04
SHC4 structural maintenance of chromosomes 4-like 1 (Yeast)	NM_004317	3.04
Consensus includes gb:BE548333 /FEA=EST /DB_XREF=gi:10579538 /DB_XREF=IMAGE:3933782 /UG=Hs.324648 Homo sapiens cDNA FLJ13700 fts, clone PLA	NM_005496	3.04
hypothetical protein FLJ10534	BE548333	3.04
FTS206 binding protein 2 (13AD)	NM_018128	3.04
arabinoside 5-epoxygenase-activating protein	NM_004470	3.04
interleukin enhancer binding factor 3, 300D	NM_001629	3.03
the GTP dissociation inhibitor (GDI) beta	AF141970	3.03
GDI1 general control of amino-acid synthesis 1-like 1 (yeast)	NM_001175	3.03
	D86973	3.03

replication factor C (activator 1) 5 (56-560)	NM_007370	2.89
polymerase (DNA-directed), delta 4	NM_021173	2.89
arc family associated phosphoprotein 2	AC03999	2.88
prosealin 1 (Nichester disease 3)	NM_007318	2.88
cytoskeleton-associated protein 1	BC005969	2.88
integrin, alpha 4 (antigen CD490, alpha 4 subunit of VLA-4 receptor)	NM_000885	2.88
NMA1228 protein	A708767	2.88
mannosidase, alpha class 2B, member 1	U68567	2.88
ADP-ribosylation factor 4	AL537042	2.87
CGA-150 protein	AF061730	2.87
hypothetical protein M6ZG555	NM_024339	2.87
protein phosphatase 2 (formerly 2A), regulatory subunit A (PR 65), beta isoform	NM_002716	2.87
interleukin 21 receptor	NM_021798	2.87
thyroid receptor interacting protein 15	NM_004236	2.87
splicing factor, arginine/serine-rich 7 (3500)	AF524053	2.87
rna homolog gene family, member H	NM_004310	2.87
SEC13-like 1 (S. cerevisiae)	NM_030673	2.86
CD4-MOT transcription complex, subunit 3	NM_014516	2.86
synovial sarcoma translocation, chromosome 18	NM_005637	2.86
NMA1513 protein	NM_017542	2.86
FES06 binding protein 1A (1200)	A936769	2.86
PRF2 periodic cryptophan protein homolog (yeast)	U56085	2.86
HSP134 protein	NM_014169	2.86
transcriptional adaptor 3 (ADA3, yeast homolog)-like (PCAF histone acetylase complex)	A024982	2.85
transferrin receptor (990, CD71)	NM_003234	2.85
ubiquitin ligase factor E48 (UBE2 homolog, yeast)	A028839	2.85
Williams-Beuren syndrome chromosome region 5	A7257135	2.85
branched chain aminotransferase 1, cytosolic	A652662	2.84
hypothetical protein	NM_015684	2.84
deklone A, fructose-bisphosphatase	A026577	2.84
maculol-like (Drosophila)	N31913	2.84
CGA-120 protein	NM_016057	2.83
selective kinase	U95339	2.83
HLA-E histocompatibility antigen, class I, G	M96886	2.83
Consensus includes gpA680302, PEA=EST /DB_XREF=gi:2556270 DB_XREF=est:ac83405.1 /CLONE=IMAGE869193 /UG=Hs.181125 immunoglobulin lambda locus	A680302	2.83
high-mobility group (nonhistone chromosomal) protein 14	NM_004965	2.83
gdy3 associated, gamma adaptin ear containing, ANF binding protein 1	NM_013365	2.83
Consensus includes gp:AL353641 /DEF=Human DNA sequence from clone RP1-158P9 on chromosome 1. Contains a putative novel gene, a laminin receptor 1 (67kD, ribosomal protein 5A)	AL353641	2.82
PEF protein with a long N-terminal hydrophobic domain (peffin)	NM_012392	2.82
mitochondrial ribosomal protein L11	NM_016050	2.82
serologically defined colon cancer antigen 16	BC001149	2.81
electronic transfer (leucoprotein, beta polypeptide	NM_001985	2.81
chromosome condensation 1	X06130	2.81
Ref1-Hirschhorn syndrome candidate 1	AF063389	2.80
SKD-domain, 6102-like, endoplatin B2	AF131863	2.80
mitochondrial ribosomal protein S11	AB049944	2.80
protein phosphatase 4 (formerly X), catalytic subunit	BC001416	2.80
A0021 protein	NM_016613	2.79
protein phosphatase 2 (formerly 2A), regulatory subunit B (PR 52), beta isoform	AA974416	2.79
similar to S. cerevisiae RER1	NM_007023	2.79
purinosis homolog 1 (Drosophila)	BC474429	2.79
heterogeneous nuclear ribonucleoprotein F	AF591354	2.79
inhibitor of growth family, member Y	AF044076	2.78
for (for/yes related) tyrosine kinase (phosphoprotein MCP94)	NM_005246	2.78
Francis aments, complementation group 6	AC064472	2.78
nuclear receptor subfamily 3, group C, member 1	U01351	2.78
immunoglobulin lambda locus	X33006	2.78
ATPase, H+-transporting, lysosomal (vacuolar protein pump) 164D	NM_001694	2.78
gdy3 transport complex 1 (90 kDa subunit)	NM_006348	2.78
hypothetical protein FL122940	NM_024571	2.77
apoptosis related protein APR-3	NM_016085	2.77
BUA0095 gene product	NM_014669	2.77

synucleophylin-like protein	A758845	2.77
G protein-coupled receptor 18	A281135	2.77
hypothetical protein PR01635	NM_019589	2.77
gpi-AF008937.1 DEF=Homo sapiens syntaxin-16c mRNA, complete cds. /FEA=mRNA PRO00=syntaxin-16c, DB_XREF=gi:2352817 /UG=Hs.102178 syntxin 16 /FL=gi:AF008937.1	AF008937	2.77
hypothetical protein FL20517	NM_017859	2.77
ribosomal protein L5	NM_009969	2.77
proteasome (prosome, macropain) subunit, beta type, 10	NM_002801	2.77
hypothetical protein NC62772	BF975273	2.77
chromosome 11 open reading frame 9	AC047770	2.76
hypothetical protein FLJ12442	NM_022908	2.76
CD164 antigen, sialomucin	BF669455	2.76
actinin 1	NM_001655	2.76
glutamic-oxaloacetic transaminase 1, soluble (aspartate aminotransferase 1)	BC004988	2.76
arginine-glutamic acid dehydrolyase (RE) repeats	AB036737	2.75
hypothetical protein FLJ14040	AK024102	2.75
chromosome 11 open reading frame 10	NM_014206	2.75
PRM4/STK19D splicing factor	U82756	2.75
transformation/transcription domain-associated protein	AC064893	2.75
BLM999 protein	AA044154	2.75
self-oxidant protein 2 (non-selenium glutathione peroxidase, acidic calcium-independent phosphoperoxidase A2)	BE869583	2.74
reticoid control protein ds3 homolog	NM_014953	2.74
DNA segment, single copy probe LUC-UU/UC-CAT (deleted in polyposis)	AA814140	2.74
bacitracin dehydrogenase 3 (NAD+) beta	AF023266	2.74
SMY3 suppressor of mit fms 3 homolog 1 (yeast)	NM_006936	2.73
translocase of outer mitochondrial membrane 20 (yeast) homolog	NM_014765	2.73
ribosomal protein L24	NM_009986	2.73
UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylglucosaminyltransferase 2 (GalNAc-T2)	NM_004481	2.73
BT33 associated nuclear protein	NM_017869	2.73
hypothetical protein FLJ12298	AK023360	2.73
ribosomal protein S10	BC004334	2.73
RNA4, member RAS oncogene family	NM_004578	2.73
QAM731 protein	BE881529	2.72
bacitracin dehydrogenase 3 (NAD+) beta	AF023265	2.72
v-cr1 melan sarcoma virus C110 oncogene homolog	NM_014647	2.72
serine (or cysteine) proteinase inhibitor, clade B (ovalbumin), member 1	NM_030666	2.72
inhibitor of DNA binding 3, dominant negative helix-loop-helix protein	NM_002167	2.72
hypothetical protein FLJ22837	NM_025165	2.72
methionine-tRNA synthetase	NM_004990	2.72
hypothetical protein FLJ22795	NM_025084	2.72
DEAD/H (Arg-Glu-Ala-Arg/His) box polypeptide 35	NM_021931	2.71
glutathione peroxidase 4 (phospholipid hydroperoxidase)	NM_002085	2.71
hypothetical protein FLJ11301	AL136897	2.71
Consensus includes gi:BP796470 /FEA=EST /DB_XREF=gi:2101524 /DB_XREF=est:60225992.6F1 /FLONE=IMAGE:4342999 /UG=Hs.284275 Homo sapiens PAN2 mRNA, complete cds. /FL=BP796470	NM_012088	2.71
poly(A) binding protein, nuclear 1	NM_004643	2.71
polymerase (RNA) II (DNA directed) polypeptide D	NM_004605	2.71
proliferation-associated 254, 3880	BF669264	2.71
QAM542 gene product	AA521233	2.71
transmethyl-RNA synthetase	NM_003191	2.71
POE3 (Lys-Arg-Glu-Leu) endoplasmic reticulum protein retention receptor 1	NM_006801	2.70
phosphatase 1	NM_002676	2.70
putative protein similar to heavy (transcriptin)	NM_005768	2.70
cytochrome P450, subfamily 51 pseudogene 2	U40033	2.70
c-est tyrosine kinase	NM_004383	2.70
ATPase, Na+/K+ transporting, beta 3 polypeptide	U51478	2.70
BU33 binding unithinized by benzimidazoles 3 homolog (yeast)	AF081496	2.69
peroxiredoxin 2	L19185	2.69
quinoid dihydropteridine reductase	BC000576	2.69
mitochondrial carrier homolog 1	AF189289	2.69
protein kinase C and casein kinase substrate in neurons 2	NM_007229	2.69
hypothetical protein FLJ11301	AL136897	2.69
putative nuclear protein	AF281030	2.69

soluble carrier family 25 (mitochondrial carrier, citrate transporter), member 1	U25147	2.58
hypothetical protein FU10769	NM_018210	2.58
TIA1 cytosolic granule-associated RNA binding protein	BF692742	2.58
cAMP responsive element modulator	NM_001881	2.58
ubiquitin specific protease 23	AA621731	2.58
hypothetical protein FU10702	NM_018184	2.58
UDP glycosyltransferase 2 family, polypeptide B15	NM_001076	2.57
polymerase (RNA) II (DNA directed) polypeptide 6	NM_002896	2.57
small glutamine-rich intratectopeptide repeat (TPR)-containing	NM_003021	2.57
CGI-56 protein	AL022316	2.57
transmembrane 7 superfamily member 2	AF086304	2.57
DNAZ756-10123 protein	BC002873	2.57
peroxylhydroxyl isomerase A (cydophilin A)	NM_021130	2.57
RAM0625 protein	NC6463	2.57
jumping translocation breakpoint	BC004239	2.56
calc-associated polypeptide, 30kD	BF247098	2.56
cell division cycle 2, G1 to S and G2 to M	088357	2.56
RAM0906 protein	AA502912	2.56
MA21 homolog (S. pombe)	BC289967	2.56
branched chain aminotransferase 2, mitochondrial	NM_001190	2.56
peroxyl-like homodomain transcription factor 1	NM_002653	2.56
hematopoietic cell-specific Lyn substrate 1	NM_005335	2.56
membrane component, chromosome 11, surface marker 1	NM_005898	2.56
anestesia resistance protein ARS2	NM_015908	2.56
cleavage and polyadenylation specific factor 1, 160kD subunit	U37012	2.56
hypothetical protein FU0244	BC002492	2.55
G1 to S phase transition 1	NM_002094	2.55
RAM0233 gene product	NM_014745	2.55
similar to S. pombe dim1+	NM_006701	2.55
MAO1 mitotic arrest deficient-like 1 (yeast)	NM_003550	2.55
adrenic phosphotransferase	NM_004485	2.55
Consensus includes g0130250, FFA=EST /B8_38EF=gi:11255432 /B8_38EF=est:G018903081 /CLONE=IMAGE:4131673 /UG=Hs.315103 ESTs. Moderately similar to DNA-binding pro	BF308250	2.55
arginase-rich, related in early stage tumors	NM_006610	2.55
interleukin 21 receptor	AF269133	2.55
clathrin, heavy polypeptide (Hc)	NM_004859	2.55
ubiquitin specific protease 24	AB028980	2.54
immunoglobulin lambda locus	AF043566	2.54
programmed cell death 8 (apoptosis-inducing factor)	NM_004208	2.54
RAE1 RNA export 1 homolog (S. pombe)	U85943	2.54
isolated (myo)-1 (or 4)-magnosphatase 2	NM_014214	2.54
protein regulator of cytokinesis 1	NM_003581	2.54
hypothetical protein FU12973	NM_024908	2.54
RAM0088 protein	W87689	2.54
NSPC023 protein	NM_014047	2.54
with 2 (leu1)	AF199015	2.54
PAI-1 mRNA-binding protein	NM_015640	2.54
RAM0635 gene product	AB022669	2.53
pseudouridylylase synthase 1	NM_025215	2.53
RAM0819 protein	AB020626	2.53
chromosome 12 open reading frame 8	NM_006817	2.53
stand regulatory element binding transcription factor 2	BE513151	2.52
Wolf-Hirschhorn syndrome candidate 1	BF111870	2.52
coxin kinase 2, alpha prime polypeptide	NM_001896	2.52
Consensus includes g014708130, FFA=EST /B8_38EF=gi:10725395 /B8_38EF=est:AV708130 /CLONE=LOC404405 /UG=Hs.283965 Homo sapiens scavenger receptor class B type III SR-I	AF043290	2.52
Pea homolog 2 (E. coli)	NM_014164	2.52
synaptic nucle expressed gene 1b	NM_018358	2.52
FXD domain-containing ion transport regulator 5	AB009358	2.51
hypothetical protein FU11198	NM_002817	2.51
mitogen-activated protein kinase kinase 7	BF663141	2.51
proteasome (prosome, macropain) 26S subunit, non-ATPase, 13	AF758755	2.51
with 2 (leu1)	NM_004781	2.50
proteasome (prosome, macropain) activator subunit 3 (PA28 gamma, Ki)		
vesicle-associated membrane protein 3 (clathrovin)		

chromosome 2 open reading frame 3	AC05034	2.50
ribosomal protein S25	AA88388	2.50
arsenite resistance protein ARS2	BE546076	2.50
a disintegrin and metalloproteinase domain 10	AU135154	2.50
RNA11228 protein	BC011224	2.50
intran light chain 1 show a	NM_002475	2.50
ATP synthase, H+ transporting, mitochondrial F0 complex, subunit g	AL050277	2.50
mitochondrial ribosomal protein L22	NM_014180	2.49
heterogeneous nuclear ribonucleoprotein H3 (H39)	AF132363	2.49
starfish, RNA binding protein (Drosophila)	AI136601	2.49
polypyridine tract binding protein (heterogeneous nuclear ribonucleoprotein I)	X52006	2.49
S-phase kinase-associated protein 2 (p45)	BC001441	2.49
estrogen-related receptor alpha	L36487	2.49
FKBP6 binding protein 1A (1260)	NM_000801	2.49
hypothetical protein DNF2p56A02822	NM_030786	2.49
regulator of telomere transcripts 1	066988	2.49
HSP162 protein	NM_014183	2.49
CXK6 cell division cycle 6 homolog (S. cerevisiae)	NM_001254	2.49
gRNU_004602.1 / DUF=Homo sapiens PK4 and SFPS1 interacting protein 2 / DUF_XREF=g-4758869 /	NM_004602	2.48
glyoxylate reductase/hydroxypyruvate reductase	AK026752	2.48
chaperonin containing TCP1, subunit 8 (beta)	NM_006595	2.48
hypothetical gene M6C15955	W67887	2.48
aldol-bis reductase family 1, member 81 (aldose reductase)	NM_001628	2.48
tyrosine 3-monooxygenase/diacylglycerol 5-monooxygenase activation protein, zeta polypeptide	U28964	2.48
Barakoid growth factor receptor 1 (fms-related tyrosine kinase 2, Phleifer syndrome)	NM_023107	2.48
low density lipoprotein receptor (familial hypercholesterolemia)	NM_000527	2.47
HSPC015 protein	NM_014026	2.47
MDZ3 homolog 8 (S. cerevisiae)	AL527855	2.47
RNA binding motif protein 14	NM_006328	2.47
C64-96 protein	AL023316	2.47
anexin A2	NM_004039	2.47
lamin B2	M94363	2.47
hypothetical protein hCLA-iso	NM_030937	2.46
lech-lea protein C3P1	NM_021633	2.46
RAM1856 protein	AK026933	2.46
MAP kinase-kinase/serine/threonine kinase 1	BC022755	2.46
chromium-activated protein kinase kinase kinase 7	AF218074	2.46
HLA-E histocompatibility antigen, class I, E	M98685	2.46
secretory carrier membrane protein 3	NM_005698	2.46
galactosidase, beta 1	NM_004404	2.46
homolog of yeast Hase MRP/RNase P protein Pop5	NM_015918	2.45
haryophorin (imporin) beta 1	AK004941	2.45
immunoglobulin lambda chain joining 3	X57812	2.45
gRNU_006326.1 / DUF=Homo sapiens seven transmembrane domain protein (NFE14), mRNA / FEA=mRNA / EN=NFE14 / PROD=seven transmembrane domain protein / DUF_XREF=g-541	NM_006326	2.45
proteasome (prosome, macropain) 26S subunit, ATPase, 1	NM_002802	2.45
hypothetical protein DNF2p4342216	AK004475	2.45
ribosomal protein S15a	NM_001019	2.44
general transcription factor II, 1	NM_001518	2.44
ubiquitin-conjugating enzyme E2D 3 (UBC475 homolog, yeast)	NM_003340	2.44
ribonuclease H1	BC534527	2.44
lipic acid synthetase	BC532462	2.44
Consensus includes gR-8332462 / FEA=EST / DUF_XREF=est-6024328371 / K1ONE=IMAGE4550505 / UG=Hs.3343 phosphoglycerate dehydrogenase	8F056778	2.44
general transcription factor IIF, polypeptide 4 (52X0 subunit)	NM_001517	2.44
hypothetical protein FLJ20360	NM_017782	2.44
RNA1028 protein	AB028551	2.44
MDH1 dehydrogenase (ubiquinone) 1 alpha subcomplex, 4 (960, MLRQ)	NM_002489	2.43
ribosomal protein L7a	NM_000972	2.43
thiosulfate and one amino acid protein kinase	AK007163	2.43
Btk finger protein Z25	AK074331	2.43
TM6 family member-associated H708 activator	NM_004180	2.43
carbamoyl-phosphate synthetase 2, aspartate transcarbamylase, and dihydroorotase	NM_004341	2.43
Tet1-Hirschhorn syndrome candidate 1	BE793789	2.43
MDL18 protein	AW248552	2.43

nucleotide binding protein	NM_025233	2.35
cell division cycle 2-like 2	AL031282	2.35
CDL18 protein	NM_015947	2.35
tumor necrosis factor receptor superfamily, member 7	NM_001242	2.34
3'-phosphoadenosine 5'-phosphatidate synthase 1	AF032026	2.34
guanine nucleotide binding protein (G protein), alpha 13	NM_006572	2.34
anaphase-1 interacting protein 1; actin receptor interacting protein 1	AC004990	2.34
hypothetical protein FLJ12806	NM_022831	2.34
CCR4-NOT transcription complex, subunit 8	AF180476	2.34
Consensus includes gi:AL365505 /DEF=human DNA sequences from clone RP11-382A12 on chromosome 20 Contains ESTs, STSs, GSSs and CpG islands. Contains the 5' part of the gene for AL365505		2.34
gi:013287.1 /DEF=human mRNA for alternative splicing product of glucocorticoidase pseudogene, complete cds. /FEA=mRNA /PROD=glucocorticoidase /DB_XREF=gi:496370 /UG=Hs.013287		2.34
casein kinase 1, delta	NM_001893	2.34
serine/threonine kinase 24 (STE20 homolog, yeast)	AA586774	2.34
pyridine-5-carboxylate synthetase (glutamate gamma-semialdehyde synthetase)	NM_002860	2.33
Consensus includes gi:AA485687 /FEA=EST /DB_XREF=est:ab381033.1 /CLONE=IMAGE:643101 /UG=Hs.284235 ESTs		2.33
lecithin:phosphatidylcholine transferase	AA486687	2.33
lecithin:phosphatidylcholine transferase	M32439	2.33
phosphatidylcholine transferase	NM_002627	2.33
DnaI (Hsp46) homolog, subfamily 8, member 6	NM_002627	2.33
heterogeneous nuclear ribonucleoprotein C (C1/C2)	AF080569	2.33
desmin (actin depolymerizing factor)	NM_004500	2.33
WW domain binding protein 11	BF697964	2.33
WW domain binding protein 11	AF118023	2.33
gi:NM_016402.1 /DEF=Homo sapiens hypothetical protein (HSPC140), mRNA. /FEA=mRNA /GEN=HSPC140 /PROD=hypothetical protein /DB_XREF=gi:7705472 /UG=Hs.279868 hypothetical		2.33
HSPC053 protein	NM_014151	2.33
malate dehydrogenase 1, MD (soluble)	NM_005917	2.33
H2A histone family, member Y	AF044286	2.33
membrane and coiled-body phosphoprotein 1	021262	2.32
Meis6 (D, melanogaster) like protein	NM_005787	2.32
gi:NM_014128.1 /DEF=Homo sapiens PR00470 protein (PR00470), mRNA. /FEA=mRNA /GEN=PR00470 protein /DB_XREF=gi:7662555 /UG=Hs.278937 PR00470 prot		2.32
hypothetical protein FLJ10083	NM_014128	2.32
DUF25600120 protein	NM_017992	2.32
hypopharynx-4RNA synthetase	NM_014077	2.32
adaptor-related protein complex 2, beta 1 subunit	NM_004184	2.32
general transcription factor IIF, polypeptide 1 (7MD subunit)	AL567295	2.32
SEC14-like 1 (S. cerevisiae)	EC000120	2.32
cytochrome b5 outer mitochondrial membrane precursor	NM_003003	2.32
D-dephospho tauroxanthone	AW235051	2.31
gi:NM_020382.1 /DEF=Homo sapiens PRSET domain containing protein 07 (SET07), mRNA. /FEA=mRNA /GEN=SET07 /PROD=PRSET domain containing protein 07 /DB_XREF=gi:9966854 /NM_020382		2.31
synthase 10	NM_001355	2.31
Consensus includes gi:H97931 /FEA=EST /DB_XREF=gi:1118816 /DB_XREF=est:tp0402.1 /CLONE=IMAGE:260738 /UG=Hs.173108 Homo sapiens cDNA: FLJ1897 fs, clone HEP03447		2.31
protein kinase, cAMP-dependent, catalytic, beta	NM_002731	2.31
peroxisomal membrane protein 2 (22B0)	NM_018663	2.31
interleukin 10 receptor, beta	BC019003	2.31
hypothetical protein FLJ10803	AF65280	2.31
Consensus includes gi:38249399 /FEA=EST /DB_XREF=gi:12759426 /DB_XREF=est:605319652F1 /CLONE=IMAGE:4414895 /UG=Hs.110771 Homo sapiens cDNA: FLJ1904 fs, clone HE		2.31
antigen identified by monoclonal antibody Kc-67	BC249599	2.31
GyrE-like protein coxapene	AU132185	2.31
hypothetical protein FLJ20257	AL542571	2.31
JAK binding protein	NM_019606	2.31
hypothetical protein FLJ11773	AB005043	2.31
mitotic spindle coiled-coil related protein	NM_021934	2.31
CDK45 cell division cycle 45-like (S. cerevisiae)	NM_006461	2.31
interferon-related developmental regulator 2	NM_003504	2.30
NOL18 protein	BC001327	2.30
nucleoside 5MD	NM_018044	2.30
tyrosyl-DNA synthetase	AF527238	2.30
diaphanous homolog 1 (Drosophila)	AW245400	2.30
thymosin, beta 10	AF051782	2.30
ubiquitin p30 (R640-containing) subunit B 1	NM_021109	2.30
cathepsin, light polypeptide (Lch)	NM_005886	2.30
cell division cycle 2-like 2	NM_001834	2.30
hypothetical protein FLJ21617	AL031282	2.30
cat eye syndrome chromosome region, candidate 5	AK025270	2.29
Consensus includes gi:M687738 /FEA=EST /DB_XREF=gi:4899032 /DB_XREF=est:tp3g08.1 /CLONE=IMAGE:2206910 /UG=Hs.699 pepitidiprosyl isomerase B (cyclophilin B)		2.29
	NM_017829	2.29
	AB87738	2.29

Consensus includes gb:AL137162 /DEF=Human DNA sequence from clone RPS-843L14 on chromosome 20. Contains ESTs, STSs and GSSs. Contains a novel gene and the 5' part of a gene to AL137162	2.29
gb:AF060511.1 /DEF=Homo sapiens clone 016s10 My016 protein mRNA. complete cds. /FEA=mRNA /PROID=My016 protein /DB_XREF=gi:12001971 /UG=Hs.161634 Homo sapiens cDNA: NM_014630	2.29
RIAA0211 gene product	2.29
isocitrate dehydrogenase 3 (NAD+) alpha	2.29
MD-1, BP105-associated	2.29
transforming growth factor, beta 2	2.29
ring finger protein 10	2.29
gb:NM_016007.1 /DEF=Homo sapiens (LOC51625), mRNA. /FEA=mRNA. KEN=LOC51625 /PROID=C61-S9 protein /DB_XREF=gi:7706297 /UG=Hs.279867 CG1-S9 protein /FI NM_016007	2.29
W3787	2.28
hypothetical protein NC02731	2.28
acetyl-Coenzyme A acyltransferase 1 (peroxisomal 3-oxoacyl-Coenzyme A thiolase)	2.28
glutathyl-prolyl-GNA synthetase	2.28
Consensus includes gb:AK000185.1 /DEF=Homo sapiens cDNA FLJ20178 fs, clone COL09990. /FEA=mRNA /DB_XREF=gi:7020104 /UG=Hs.306389 Homo sapiens cDNA FLJ20178 fs, clone AK000185	2.28
hypothetical protein FLJ20764	2.28
protease, serine, 25	2.28
mitogen-activated protein kinase kinase 12	2.28
RCS1-like (yeast)	2.28
regulator of G-protein signalling 10	2.28
adrenocortical transition termination factor 1	2.28
ubiquitin-cytochrome c reductase hinge protein	2.28
scaffold attachment factor 8	2.28
hypothetical protein MG2840 similar to a putative glucosyltransferase	2.28
calumenin	2.27
sialyltransferase 7D ((alpha-N-acetylneuraminyl-2,3-beta-galactosyl-1,3)-N-acetyl galactosaminide alpha-2,6-sialyltransferase)	2.27
potassium channel, subfamily K, member 1 (TWIK-1)	2.27
Rho GDP dissociation inhibitor (GDI) alpha	2.27
RNA binding motif protein, X chromosome	2.27
defender against cell death 1	2.27
mG2 homolog 3 (E. coli)	2.27
polymerase (RNA) III (DNA directed) (6240)	2.27
ubiquitin A-S2 residue ribosomal protein fusion product 1	2.27
tubulin-specific chaperone d	2.27
LIM domain only 2 (Hombobin-like 1)	2.26
clathrin, light polypeptide (Lc)	2.26
protein kinase C beta 1	2.26
putative dimethyladenosine transferase	2.26
chromosome 6 open reading frame 11	2.26
protein phosphatase 6, catalytic subunit	2.26
phosphoribosylglycylamide formyltransferase, phosphoribosylglycylamide synthetase, phosphoribosylaminimidazole synthetase	2.25
hypothetical protein MG2861	2.25
Consensus includes gb:AL096428 /DEF=Human DNA sequence from clone RPS-963E22 on chromosome 20. Contains the 3' end of a novel gene similar to NY-REN-2 Antigen, 5' Cdc islands, E. AL096428	2.25
RIAA0071 protein	2.25
dCMP deaminase	2.25
adenylylase kinase 2	2.25
methylenetetrahydrofolate synthetase 2 (beta)	2.25
hypothetical protein	2.25
M-phase phosphoprotein 9	2.25
RIAA0077 protein	2.25
activity-dependent neuroprotector	2.25
serine protease inhibitor, Kunitz type, 2	2.25
clathrin, light polypeptide (Lc)	2.24
Consensus includes gb:AF062978 /FEA=EST /DB_XREF=gi:5755591 /DB_XREF=est:W2505.x1 /CLONE=IMAGE250513 /UG=Hs.288908 WAS protein family, member 2 /FI=gb:NM_00659 /AP062978	2.24
interleukin 4 receptor	2.24
fragile X mental retardation, autosomal homolog 1	2.24
CD48 antigen (B-cell membrane protein)	2.24
SMC4 structural maintenance of chromosomes 4-like 1 (yeast)	2.24
Eti-domain containing 1	2.24
aryl hydrocarbon receptor interacting protein	2.24
POU domain, class 2, associating factor 1	2.24
ECMT	2.24
Consensus includes gb:AU145293 /FEA=EST /DB_XREF=gi:11006814 /DB_XREF=est:AU145293 /CLONE=HEHBA1004405 /UG=Hs.61773 Homo sapiens cDNA FLJ11648 fs, clone HEHBA1 AU145293	2.24
phosphoglycerate dehydrogenase	2.24
calreticulin	2.23